Disclosure; SEQ ID NO 8; 135pp; English.

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The invention relates to a method of identifying a binding partner for synaptic vesicle protein 2 (SV2), involves incubating SV2 protein with levetiracetam and a potential binding partner and determining if the potential binding partner modulates the binding of levetiracetam to the SV2 protein or fragment, thus identifying a binding partner for the protein. The method is useful for identifying a binding partner for synaptic vesicle protein 2 and the identified binding partner for for tradting epilepsy, epileptogenesis, seizure disorders, convulsions, depression, anxiety, cerebral ischaemia, myotonia, stroke, Tourette's syndicme, neonatal cerebral haemorrhage, Parkinson's disease, Alzheimer's
                            Identifying binding partner for synaptic vesicle protein 2 (SV2) for treating epilepsy, Parkinson's disease, Alzheimer's disease, involves determining binding partner which modulates binding of levetiracetam or its analog to SV2 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease and dementia. The present sequence represents the amino acid sequence of the human synaptic vesicle protein 2, SVOP.
                                                                                                                                                          Disclosure; SEQ ID NO 8; 63pp; English.
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Sequence 548 AA;

Gaps ; 30.0%; Score 6; DB 8; Length 548; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels 6; Conservative Query Match Best Local Similarity Matches ð

o;

8 LGWRWL 13

235 LGWRWL 240

RESULT 42 ADP44588 

ADP44588 standard; protein; 548 AA (first entry) 09-SEP-2004 ADP44588;

Human synaptic vesicle protein OP (SVOP).

relaxant, haemostatic, neuroprotective; antiparkinsonian; nootropic; endocrine; antidiabetic; seizure; epilepsy; Parkinson's; Alzheimer's; cognitive; movement; bipolar; neurodegenerative; dementia; gigantism; dwarfism; adrenal-medulla; hypoglycaemia; circulation shock; human; SVOP. neurological disorder; synaptic vesicle function; endocrinopathy; hormonal; SV2; synaptic vesicle protein 2; anticonvulsant; neuroleptic; antimanic; antidepressant; tranquiliser; antimigraine; analgesic; cerebroprotective; vasotropic; vulnerary; muscle relaxant; acaricide;

Homo sapiens.

WO2004051222-A2

02-DEC-2003; 2003WO-US038122.

03-DEC-2002; 2002US-0430372P. 30-SEP-2003; 2003US-0506764P.

(UNIO ) UCB SA.

Fuks Lynch B, Nocka K,

WPI; 2004-461170/43. N-PSDB; ADP44587. Treating neurological disorder associated with synaptic vesicle function such as seizure, epilepsy, endocrinopathy or hormonal diseases, involves administering compound that modulates function or activity of SV2 proteins

The invention relates to a novel method for treating a neurological disorder associated with synaptic vesicle function, endocrinopathy or hormonal disease which comprises administering a compound or agent that condulates a function or activity of an SV2 (synaptic vesicle protein.) Protein. The method of the invention has anticonvulsant, neuroleptic, or antimanic, antidepressant, tranquiliser, antimagraine, analgesic, cerebroprotective, vasotropic, vulnerary, muscle relaxant, nacmotatic, neuroprotective, antimagraine, analgesic, cerebroprotective, vasotropic, vulnerary, muscle relaxant, notropic, condocrine and antidabetic applications. The method may be useful for treating a neurological disorder, endocrinopathy or hormonal disease associated with synaptic vesicle function. The neurological disorder is chosen from selutery, parkinson's disease, Alzheimer's disease, cognitive disease and dementia. The endocrinological disorder is neurodegenerative disease and dementia. The endocrinological disorder is chosen from endocrinopathies involving hypersecretion or hyposecretion of at least one hormone, gigantism, dwarfism, adrenal-medulla-related that of the human synaptic vesicle protein OP (SVOP) of the invention. 0; Gaps ., 30.0%; Score 6; DB 8; Length 548; 100.0%; Pred. No. 2.8e+02; Live 0; Mismatches 0; Indels 6; Conservative 235 LGWRWL 240 Local Similarity 8 LGWRWL 13 Sequence 548 AA; Query Match Matches RESULT 43 à g

ADP44596 standard; protein; 548 AA. ADP44596

(first entry) 09-SEP-2004 

Norway rat synaptic vesicle protein OP (SVOP).

neurological disorder; synaptic vesicle function; endocrinopathy; hormonal; SV2; synaptic vesicle protein 2; anticonvulsant; neuroleptic; antimanic; antidepressant; tranquiliser; antimigraine; analgesic; cerebroprotective; vasotropic; vulnerary; muscle relaxant; acaricide; relaxant; haemostatic; neuroprotective; antiparkinsonian; nootropic; endocrine; antidiabetic; seizure; epilepsy; Parkinson; s. Alzheimer's; cognitive; movement; bipolar; neurodegenerative; dementia; gigantism; dwarfism; adrenal-medulla; hypoglycaemia; circulation shock; Norway rat;

WO2004051222-A2.

17-JUN-2004.

02-DEC-2003; 2003WO-US038122.

03-DEC-2002; 2002US-0430372P 30-SEP-2003; 2003US-0506764P

(UNIO ) UCB SA.

Fuks Lynch B, Nocka K,

WPI; 2004-461170/43.

N-PSDB; ADP44595

Treating neurological disorder associated with synaptic vesicle function such as seizure, epilepsy, endocrinopathy or hormonal diseases, involves

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RESULT 45
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                                                           The invention relates to a novel method for treating a neurological disorder associated with synaptic vesicle function, endocrinopathy or disorder associated with synaptic vesicle function, endocrinopathy or hormonal disease which comprises administaring a compound or agent that modulates a function or activity of an SV2 (synaptic vesicle protein 2) protein. The method of the invention has anticonvulsant, neuroleptic, antimanic, antidepressant, tranquiliser, antimigraine, analgesic, cerebroprotective, vasotropic, vulnerary, wescle relaxant, acaricide, relaxant, haemostatic, neuroprotective, antiparkinsonian, nootropic, relaxant, haemostatic, neuroprotective, antiparkinsonian, nootropic, endocrine and antidiabetic applications. The method may be useful for treating a neurological disorder, endocrinopathy or hormonal disease, cospective disorders, movement disorders, pipolar disorder is cognitive disorders, movement disorders, bipolar disorders, neurodegenerative disease and dementia. The endocrinological disorder is chosen from endocrinopathies involving hypersecretion or hyposecretion of at least one hormone, gigantism, dwarfism, adrenal-medulla-related diseases, hypoglycaemia and circulation shock. The current sequence is continuation should be an endocrinopathy content of the current sequence is that of the Norway rat synaptic vesicle protein Op (SVOP) of the
                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enantiomeric conversion; amine; homochiral; enzyme; oxidation catalysis; stereoselective; microbial; monoamine oxidase; MAO; enantioselectivity; deracemization; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             348
/note= "The wild-type residue of Met is substituted with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note≈ "The wild-type residue of Asn is substituted with
                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aspergillus niger monoamine oxidase protein variant N336S + M348K.
administering compound that modulates function or activity of SV2
                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "All Xaa residues signify a stop codon"
                                                                                                                                                                                                                                                                                                                                                     30.0%; Score 6; DB 8; Length 548; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                        Disclosure, SEQ ID NO 16, 135pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ney Location/Qualifiers Misc-difference j. .610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD15932 standard; protein; 610 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                   6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    235 LGWRWL 240
                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                           8 LGWRWL 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aspergillus niger.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                  Sequence 548 AA;
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                                                                                                                                                                                                                                                                                         that of th
invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD15932;
                                                                                                                                                                                                                                                                                                                                                          Query Match
                     proteins
                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD15932
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enantiomeric conversion; amine; homochiral; enzyme; oxidation catalysis; stereoselective; microbial; monoamine oxidase; MAO; enantioselectivity; deracemization; mutant; mutein.
                                                                                                                                                                                                                                                                                                                               The invention relates to a novel method for the enantiomeric conversion of amines. The method comprises treating a homochiral amine or a mixture of amine enantioners with an enzyme capable of catalyshing oxidation of the amine in a stereoselective manner and, subsequently or simultaneously, treating with a reducing agent. A further method of the invention is useful for directing the evolution of an amine oxidation enzyme e.g., microbial monoamine oxidase (MAO), preferably Aspergillus niger monoamine oxidase or its variant. This method is also useful for increasing the enantioselectivity of an amine oxidase enzyme. The enantioselective amine oxidase enzyme of the invention is useful in deracemization of amines This sequence represents a variant Aspergillus niger monoamine oxidase protein of the invention.
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                                                                                                               Enantiomeric conversion of amines comprises treating a homochiral amine or mixture of amine enantiomers with enzyme capable of catalyzing oxidation of amine and subsequently or simultaneously treating with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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'note= "All Xaa residues signify a stop codon"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30.0%; Score 6; DB 7; Length 610; 100.0%; Pred. No. 3e+02; cive 0; Mismatches 0; Indels
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Mahmoudian M;
   Turner NJ,
                                                                                                                                                                                                                                                                                    Claim 26; SEQ ID NO 3; 43pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Conservative
   Alexeeva MV, Enright A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GLAX ) GLAXO GROUP LTD
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                                                              WPI; 2003-779267/73
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 WALGWR 11
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Aspergillus niger.
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                                                                                                                                                                                                                                   reducing
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Enantiomeric conversion of amines comprises treating a homochiral amine or mixture of amine enantiomers with enzyme capable of catalyzing oxidation of amine and subsequently or simultaneously treating with a reducing agent

Claim 24; SEQ ID NO 2; 43pp; English

The invention relates to a novel method for the enantiomeric conversion of amines. The method comprises treating a homochiral amine or a mixture of amine enantiomers with an enzyme capable of catalyshing oxidation of the amine in a stereoselective manner and, subsequently or simultaneously, treating with a reducing agent. A further method of the invention is useful for directing the evolution of an amine oxidation enzyme e.g., microbial monoamine oxidase (MAO), preferably Aspergillus niger monoamine oxidase or its variant. This method is also useful for increasing the enantioselectivity of an amine oxidase enzyme. The enantioselective amine oxidase enzyme of the invention is useful in deracemization of amines. This sequence represents a variant Aspergillus niger monoamine oxidase protein of the invention.

Sequence 610 AA;

Gaps . 0 Length 610; 0; Indels 30.0%; Score 6; DB 7; I 100.0%; Pred. No. 3e+02; 0; Mismatches 6; Conservative Best Local Similarity Query Match Matches ð

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6 WALGWR 11

459 WALGWR 464

ADD15930

ADD15930 standard; protein; 610 AA. ADD15930;

(first entry) 15-JAN-2004 Wild-type Aspergillus niger monoamine oxidase protein.

enantiomeric conversion, amine; homochiral; enzyme; oxidation catalysis; stereoselective; microbial; monoamine oxidase; MAO; enantioselectivity; deracemization; wild-type.

Aspergillus niger.

Location/Qualifiers .610 Misc-difference 1.

/note= "All Xaa residues signify a stop codon"

WO2003080855-A2

02-OCT-2003

19-MAR-2003; 2003WO-GB001198

19-MAR-2002; 2002GB-00006415

(GLAX ) GLAXO GROUP LTD.

Mahmoudian M; Turner NJ, Enright A, Alexeeva MV,

WPI; 2003-779267/73

Enantiomeric conversion of amines comprises treating a homochiral amine or mixture of amine enantiomers with enzyme capable of catalyzing amine and subsequently or simultaneously treating with or mixture of amine enantiomers with enzyme oxidation of amine and subsequently or aimin agent reducing

Example 7; SEQ ID NO 1; 43pp; English.

The invention relates to a novel method for the enantiomeric conversion

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of amines. The method comprises treating a homochiral amine or a mixture of amine enantioners with an enzyme capable of catalysing oxidation of the amine in a stereoselective manner and subsequently orthat or simultaneously, treating with a reducing agent. A further method of the invention is useful for directing the evolution of an amine oxidation enzyme e.g., microbial monoamine oxidase (MAO), preferably Aspergillus niger monoamine oxidase or its variant. This method is also useful for increasing the enantioselectivity of an amine oxidase enzyme. The denatioselective amine oxidase enzyme of the invention is useful in deracemization of amines. This sequence represents the wild type Aspergillus niger monoamine oxidase protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polynucleotides derived from Coryneform bacteria, for identifying mutation point of a gene, measuring expression of a news and notice.
                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yokoi H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mutation point of a gene, measuring expression of a gene, analyzing expression profile or pattern of a gene and identifying homologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides a number of nucleotide and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coryneform bacterium; amino acid synthesis; vitamin; saccharide;
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                                                                                                                                                                                                                                                                                   30.0%; Score 6; DB 7; Length 610; 100.0%; Pred. No. 3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       H, Ando S, Hayashi M,
Ikeda M, Ozaki A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C glutamicum protein fragment SEQ ID NO: 3935.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAG90181 standard; protein; 755 AA.
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2000JP-00159162.
2000JP-00280988.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Corynebacterium glutamicum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organic acid synthesis.
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                                                                                                                                                                                                                                                                                                                                                                                                             459 WALGWR 464
                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                     6 WALGWR 11
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                                                                                                                                                                                                                                               Sequence 610 AA;
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Tateishi N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAG90181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 47
                                                                                                                                                                                                                                                                                                                             Matches
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Sequence 755 AA;

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This invention describes novel polymucleotides that encode membrane synthase and membrane transport proteins from Corynebacterium glutamicum. The polymucleotides are isolated from a nucleic acid library of C. glutamicum then mutated at the specially Corynebacterium glutamicum. Use that express the polymucleotides are used for containing vectors that express the polymucleotides are used for production of fine chemicals, preferably amino acids and specifically lyaine, but more generally nucleotides, nucleosides, lipids, fatty acids, diols, carbohydrates, aromatic compounds, vitamins, co-factors and enzymes. These are used in the food, animal feed, cosmetics and pharmaceutical industries. The polymucleotides, optionally as primers and probes, can also be used for identification and classification of C. probes, can also be used for identification and classification of C. glutamicum and related species, e.g. for diagnosis, for genomic mapping, functional or evolutionary studies, gene manipulation and modulation of metabolic activity. Cells that containing the polymucleotides of the invention may produce fine chemicals in better yields, with higher productivity and/or more efficiently. NOTE: This sequence is not represented in the printed specification but is available in electronic formations.
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    G. glutamicum membrane transport and synthesis-associated protein #39.

                                                                                                                                                                                                                                                                                                                                                                                                                                                         fine chemical production; lysine production; nucleotide; nucleoside; lipid; fatty acid; diol; carbohydrate; aromatic compound; vitamin; co-factor; enzyme; food; animal feed; cosmetic; pharmaceutical; gene;
                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid encoding variant forms of membrane construction and transport proteins, useful for production of fine chemicals from microorganisms, e.g. nucleotides, lipids and especially lysine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pompejus M, Schroeder H, Kroeger B, Klopprogge C;
                                                                                                                    0;
electronic format directly from the European Patent Office
                                                                            Length 755;
                                                                                                                    0; Indels
                                                                            DB 4; Ler
                                                                          30.0%; Score 6; DB 4
100.0%; Pred. No. 3.6
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     membrane synthesis; membrane transport.
                                                                                                                                                                                                                                                                                                ADL65505 standard; protein; 755 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-2001; 2001DE-01054179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-NOV-2001; 2001DE-01054179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page; 20pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corynebacterium glutamicum.
                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                    6; Conservative
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                                                                                                                                                                                                WSLWAL 195
                                                                            Query Match
Best Local Similarity
                                                                                                                                                           WSLWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB: ADL65504.
                                     Sequence 755 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DE10154179-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Haberhauer G;
                                                                                                                                                                                                                                                                                                                                                                               20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                       ADL65505;
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                                                                                                                                                                                                                                                          RESULT 48
ADL65505
                                                                                                                    Matches
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WO2003040292

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The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them The sequences are useful in diagnosis and planucleotides encoding them The sequences are useful in diagnosis and theraphological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, of for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant or production of P. aeruginosa caused components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences of Pseudomonas species using biochip technology. Sequences ABO67826-ABO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but, was obtained in electronic format from USPTO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ·;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                            Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                     0;
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 Length 755;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.0%; Score 6; DB 7; Length 800; 100.0%; Pred. No. 3.7e+02; cive 0; Mismatches 0; Indels
                                   0; Indels
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 DB 7; Ler
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 24635; 455pp; English.
30.0%; Score 6; DB 7
100.0%; Pred. No. 3.6
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rubenfield MJ, Nolling J, Deloughery C,
                                                                                                                                                                                                                                                                                            Pseudomonas aeruginosa polypeptide #8064.
                                                                                                                                                                                         ABO75889 standard; protein; 800 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              segdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00252991.
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98US-0094190P.
                                                                                                                                                                                                                                                         29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 10v.
                                      Conservative
                                                                                                                                                                                                                                                                                                                                                             Pseudomonas aeruginosa.
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                                                                                                      190 WSLWAL 195
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   Query Match
Best Local Similarity
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                                                                    3 WSLWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; ABD09460.
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                                                                                                                                                                                                                                                                                                                                                                                               US6551795-B1.
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                                                                                                                                                                                                                         AB075889;
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                                      Matches
                                                                                                                                                       RESULT 49
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AAM24250 standard; protein; 896 AA.

AAM24250;

WLRRYG 462 12 WLRRYG 17

457

g 8

RESULT 51 AAM24250

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The present invention describes a predictor set comprising a plurality of polymucleotides or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein tyrosine kinase activity or members of the protein tyrosine kinase compared to the activity of cells, comprising obtaining a sample of cells, pathway. Also described: (1) predicting whether a compound is capable of modulating the activity of cells, comprising obtaining a sample of cells, correlating the extression of the markers to the compound's ability to correlating the extression of the markers to the compound's ability to correlate what compound sensitivity or resistance of cells associated with a disease state, and (3) identifying polymucleotides and polypeptides whose expression levels correlate with a disease state, and (3) identifying polymucleotides and polypeptides that predict compound sensitivity or resistance of cells associated with a disease state, compounds, analysing the expression pattern of a microarray of polymucleotides that predict the sensitivity or resistance of cells associated with a disease state, opplyapptides, and selecting to polymucleotides or polypeptides that predict the sensitivity or resistance of cells associated with a disease state by using the expression pattern of the microarray. The polymucleotides and cervistes of cells associated with a disease state by using the corpounds that interact with protein tyrosine kinase pathways. These may be used in determining drug sensitivity in patients to allow the development of individualized cancer; based on patient response at a molecular level. The present corporation cancer; based on patient response at a molecular level. The present
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                                                                                                                                                                                                     predictor set; protein tyrosine kinase activity modulator; protein tyrosine kinase pathway; protein tyrosine kinase; cytostatic; gene therapy; drug sensitivity; genetic profile; cancer; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways.
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                                                                                                                                                                   Human src biomarker polypeptide SEQ ID NO:339.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shaw P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; SEQ ID NO 339; 139pp; English
                                        ADD14150 standard; protein; 880 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lee FY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BRIM ) BRISTOL-MYERS SQUIBB CO
                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JAN-2003; 2003WO-US001981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JAN-2002; 2002US-0350061P
                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Huang F, Fairchild CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-636735/60.
N-PSDB; ADD14752.
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                                                                                                                                                                                                                                                                                                                                                WO2003062395-A2.
                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                            01-JAN-2004
                                                                                    ADD14150;
RESULT 50
                      ADD14150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (BSTs) from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a protein of the invention
                                                                                   Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; horse;
tomato; monkey; dog; sea urchin; expressed sequence tag; EST;
diagnostics; forensic test; gene mapping; genetic disorder; biodiversity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated polypeptide for treatment of diseases, diagnostics, raising antibodies and research use.
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                                                                                                                                                                                                                                                                                                                                                                   Chen R,
                                                                                                                                                                                                                                                                                                                                                                  ou P, Qian XB, Wang Z,
Zhang J, Werhman T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 20; Page 1166-1168; 1275pp; English.
                                                             Human EST encoded protein SEQ ID NO: 1775.
                                                                                                                                                                                                                                                                                                                                                                      Zhou P, Qian XB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB79887 standard; protein; 984 AA.
                                                                                                                                                                                                                                            25-JAN-2001; 2001WO-US002687.
                                                                                                                                                                                                                                                                                   17-JUL-2000; 2000US-00617746.
03-AUG-2000; 2000US-00631451.
15-SEP-2000; 2000US-00663870.
                                                                                                                                                                                                                                                                       25-JAN-2000; 2000US-00491404
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                                      (first entry)
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nes 6; Conservative
                                                                                                                                 gene therapy; nutrition.
                                                                                                                                                                                                                                                                                                                                                               Tang YT, Liu U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              687 WLRRYG 692
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                                                                                                                                                                                                                                                                                                                                            (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 896 AA;
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                                                                                                                                                                                        WO200154477-A2.
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                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                  02-AUG-2001.
                                     12-OCT-2001
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ID AAB7
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AC AAB7
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DT 30-1
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pyrimidine bases.
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                                                         N-PSDB; AAF72006
                                                                                                                                                                            Sequence 984 AA;
                        (BADI ) BASF AG.
                                                                                                                                                                                                                                                                                                                                                                           WO200100843-A2
                                                                                                                                                                                                                                                                                                                                                                                                      23-JUN-2000;
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                                                                                                                                                                                                                                                                         fine chemical production, microorganism, organic acid, nucleoside, nonproteinogenic amino acid, purine base, pyrimidine base, nucleotide, lipid; saturated fatty acid, unsaturated fatty acid, diol, vitamin, carbohydrate, aromatic compound; cofactor; polyketide, enzyme.
                   Corynebacterium glutamicum; metabolic pathway protein; MP protein;
     Corynebacterium glutamicum MP protein sequence SEQ ID NO:508.
                                                                                                                                                                                                990E-01031453

990E-01031457

990E-01031410

990E-01031541

990E-01031541

990E-01031541

990E-01031532

990E-01031252

990E-01032125

990E-01032126

990E-01032126

990E-01032206

990E-01032208

990E-01032208

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990E-01032208
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99DE-01030476.
99DE-01031418.
99DE-01031419.
99DE-01031420.
99DE-01031428.
99DE-01031428.
99DE-01031428.
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99DE-01042077
99DE-01042079
99DE-01042087
99DE-01042088
99DE-01042088
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99DE-01033005,
99DE-01033006,
99US-0148613P.
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99DE-01040832.
99DE-01041378.
99DE-01041379.
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99DE-01041394.
99DE-01041396.
                                                                                                   23-JUN-2000; 2000WO-IB000923
                                                                                                                                                                                                                                                                                                                                                                                     99DE-01040764
                                                                                                                                                                                                                                                                                                                                                                                            99DE-01040765
                                                            Corynebacterium glutamicum.
                                                                         WO200100843-A2
                                                                                                                                                                                                             08-JUL-1999;
08-JUL-1999;
08-JUL-1999;
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09-JUL-1999;
09-JUL-1999;
09-JUL-1999;
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08-JUL-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF71753 to AAF72330 encode the Corynebacterium glutamicum metabolic pathway (MP) proteins given in AAB79634 to AAB80211. The C. glutamicum MP nucleic acids are useful for the production of fine chemicals in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             microorganisms, including organic acids, nonproteinogenic amino acids, purine and pyrimidine bases, nucleosides, nucleotides, lipids, saturated and unsaturated fatty acids, diols, carbohydrates, aromatic compounds, vitamins, cofactors, polyketides and enzymes
                                                                                                                                                                                                                                                             Nucleic acids from Corynebacterium glutamicum encoding metabolic pathway proteins, useful for producing fine chemicals in microorganisms, including organic acids, nonproteinogenic amino acids, and purine and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                     Zelder O, Haberhauer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Corynebacterium glutamicum MP protein sequence SEQ ID NO:594.
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100.0%; Pred. No. 4.4e-
                                                                                                                                                                                                                                                                                                                                                                                             Claim 20; Page 887-890; 1737pp; English.
                                                                                                                                     Schroeder H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB79930 standard; protein; 984 AA
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99US-0142101P
99US-01031415
99DE-01031419
99DE-01031420
99DE-01031424
99DE-01031434
99DE-01031443.
03-SEP-1999; 99DE-01042129.
09-MAR-2000; 2000US-0187970P.
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                                                                                                                                  Pompejus M, Kroeger B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Conservative
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Best Local Similarity
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4 SIWALG 9

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AAF71753 to AAF72330 encode the Corynebacterium glutamicum metabolic pathway (MP) proteins given in AAB79634 to AAB80211. The C. glutamicum MP nucleic acids are useful for the production of fine chemicals in microorganisms, including organic acids, nonproteinogenic amino acids, purine and pyrimidine bases, nucleosides, nucleotides, lipids, saturated and unsaturated fatty acids, diols, carbohydrates, aromatic compounds, vitamins, cofactors, polyketides and enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids from Corynebacterium glutamicum encoding metabolic pathway proteins, useful for producing fine chemicals in microorganisms, including organic acids, nonproteinogenic amino acids, and purine and pyrimidine bases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Haberhauer G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30.0%; Score 6; DB 4; Length 984; 100.0%; Pred. No. 4.4e+02; vative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 20; Page 1000-1003; 1737pp; English.
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99DE-01041396.
99DE-01042076.
99DE-01042077.
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99DE-01042086.
99DE-01042087.
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99DE-01032229.
99DE-01032230.
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99DE-01031573.
99DE-01031592.
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99DE-01032130.
99DE-01032186.
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99DE-01032926.
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99DE-01040765.
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                                                                              99DE-01031632,
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 99DE-01031457
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-137957/14.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; AAF72049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BADI ) BASF AG.
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08.7UL-1999;
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09-7UL-1999;
09-7UL-1999;
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03-SEP-1999
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27-AUG-19
31-AUG-19
31-AUG-19
31-AUG-19
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27-AUG-1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polynucleotides derived from Coryneform bacteria, for identifying mutation point of a gene, measuring expression of a gene, analyzing expression profile or pattern of a gene and identifying homologous gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mizoguchi H, Ando S, Hayashi M, Ochiai K, Yokoi H;
Senoh A, Ikeda M, Ozaki A;
                                                                                                                                                                                                        amino acid synthesis; vitamin; saccharide;
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4.7e+02;
0;
                                                                                                                                                                        C glutamicum protein fragment SEQ ID NO: 6914.
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100.0%; Pred. No. 4.7
tive 0; Mismatches
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                                                                             AAG93160 standard; protein; 1083 AA
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07-APR-2000; 2000JP-00159162.
03-AUG-2000; 2000JP-00280988.
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                                                                                                                                                                                                                                                                                                                                                         18-DEC-2000; 2000EP-00127688
                                                                                                                                                                                                                                                           Corynebacterium glutamicum
                                                                                                                                               26-SEP-2001 (first entry)
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                                                                                                                                                                                                            Coryneform bacterium; a organic acid synthesis.
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SLWALG 409
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Tateishi N,
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Watanabe
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              9805-0102487P

9805-010257P

9805-0102565P

9805-0102565P

9805-0102328B

9805-0103449P

9805-01034328P

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9805-01034328P

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9805-01034328P

9805-01034328P

9805-01034328P

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9805-0105681P

9805-0105681P

9805-0105681P

9805-0105632P

9805-0106632P

9805-0106932P

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N-PSDB; AAA37051.
    30-SEP-1998,
30-SEP-1998,
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10-OCT-1998,
01-OCT-1998,
06-OCT-1998,
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07-OCT-1998,
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27-0CT-1998;
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     Human; PRO polypeptide, membrane bound protein; receptor; diagnosis; transmembrane; secretion; immunoadhesion; pharmaceutical; screening.
                                     Human PRO1249 (UNQ632) amino acid sequence SEQ ID NO:102
                                                                                                                                                                    98US-0098716P.
98US-0098873P.
98US-0098803P.
98US-0098831P.
98US-0099831P.
98US-0099536P.
98US-0099536P.
98US-0099763P.
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98US-0099764P.
98US-0099764P.
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98US-0100684P.
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98US-0100684P.
98US-0100711P.
98US-0100711P.
98US-0100711P.
98US-010071P.
98US-010071P.
98US-010071P.
98US-010147P.
98US-0101738P.
98US-0101738P.
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                  08-AUG-2000 (first entry)
                                                                                                           WO200012708-A2
                                                                                        Homo sapiens.
                                                                                                                                                                    01-SEP-1998,
01-SEP-1998,
02-SEP-1998,
02-SEP-1998,
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16-SEP-1998,
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17-SEP-1998
AAY99369;
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Wood WI;

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receptor or secreted PRO polypeptides given in AAY99340 to AAY99462. The transmembrane and receptor PRO proteins can be used for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interactions. The polypeptides and nucleotide sequences encoding then have various industrial applications, including uses as pharmaceutical and diagnostic agents. AAA37145 to AAA3730 represent PCR primers and hybridisation probes used in the isolation of the PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Secreted and transmembrane proteins and nucleic acids designated PRO, useful as hybridization probes, in chromosome and gene mapping and gene
    New mammalian DNA sequences encoding transmembrane, receptor or secreted PRO polypeptides, useful for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interactions.
                                                                                        to AAA37144 encode the new isolated human transmembrane,
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Gurney AL, <sup>1</sup>
'A, Tumas D,
                                                                                                                                                                                                                                                           Length 1089;
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Gao W, Goddard A, Godowski PJ, Grimaldi CJ, Gu
Pan J, Paoni NF, Roy MA, Smith V, Stewart TA,
                                                                                                                                                                                                                                                          30.0%; Score 6; DB 3; Ler 100.0%; Pred. No. 4.8e+02; ive 0; Mismatches 0;
                                                                                                                                                                                                       polypeptides from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Secreted; transmembrane; gene therapy
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                                                                Claim 12; Fig 60; 773pp; English
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99US-0144758P.
99US-0145698P.
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2000WO-US000219
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                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                              687 WLRRYG 692
                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                  12 WLRRYG 17
                                                                                                                                                                                                                                    Sequence 1089 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified.
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29-OCT-1999;
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Fong S; Hillan KJ; Watanabe CK;

Claim 1; Fig 60; 787pp; English

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 The present invention relates to secreted and transmembrane proteins. These proteins and the DNA encoding them may be used as hybridization probes, in chromosome and gene mapping and in the generation of antisense RNA and DNA. They may also be used used to generate either transgenic animals or knockout animals which are in turn useful for development and screening of therapeutically useful reagents. The nucleic acids may also be used in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                          PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep; dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha; blood; chondrocyte cell; cell proliferation; cell differentiation; colon; adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.
                                                                                                                                                                      Gaps
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                                                                                                                                         30.0%; Score 6; DB 4; Length 1089; 100.0%; Pred. No. 4.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide sequence #133.
                                                                                                                                                                                                                                                                                                   AAU29156 standard; protein; 1089 AA
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14-MAR-2000; 2000US-0189320P.
15-MAR-2000; 2000US-0189320P.
15-MAR-2000; 2000US-0190828P.
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04-APR-2000; 2000US-0194449P.
04-APR-2000; 2000US-0194647P.
11-ARR-2000; 2000US-0195975P.
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2000US-0196187P.
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2000US-0198585P.
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02-MAR-2000; 2000WO-US005841.
03-MAR-2000; 2000US-0187202P-
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                                                                                                                                                                           Conservative
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                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                     Sequence 1089 AA;
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11-APR-2000;
11-APR-2000;
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98US-0088217P.
98US-0088655P.
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28-0CT-1997
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31-0CT-1997
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12-DEC-1997;
17-DEC-1997;
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01-APR-1998;
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09-JUN-1998;
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08-APR-1998;
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07-MAY-1998;
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21-APR-1998
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04-JUN-1998
        Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.

The PRO polypeptides and their associated nucleic acids can be used to detect the presence of a tumour in a mammal by comparing the level of expression of a PRO polypeptide in a test sample of cells from the animal early a control sample of normal cells, whereby a higher level of and a control sample of normal cells, whereby a higher level of expression in the test sample indicates the presence of a tumour in the mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats and rabbits but are preferably human. The polypeptides can be used to stimulate tumour necrosis factor (TNF) alpha release from human blood, the proliferation or differentiation of fondrocyte cells. The proliferation of determine the presence of tumours and also susceptibility to tumour development, particularly afternal, unng, colon, breast, prostate, rectal, cervical, or liver tumours, in mammalian cervical, cervical, or liver tumours, in mammalian cells subjects. The oligonucleotide probes specific for the PRO nucleic acids can be used for genetic analysis of individuals with genetic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                             Novel nucleic acids encoding PRO polypeptides, used to diagnose the presence of tumors, such as prostate and breast tumors, in mammals and to screen for modulators of the compounds.
                                                                                                                                             , Goddard A, Godowski PJ, Gurney AL; Wood WI, Zhang Z;
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dog, cat, cow, horse, sheep, pig, goat, rabbit, ADEPT,
antibody-dependent enzyme mediated prodrug therapy.
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                                                                                                                                            P, Chen J, Desnoyers L, Smith V, Watanabe CK, W
                                   2000WO-US020710
2000US-00644848
2000WO-US023328
2000WO-US030952
2000WO-US032678
            2000WO-US015264
2000US-0209832P
                                                                                             2000WO-US034956
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Matches 6; Conserv
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30-MAY-2000; 2
02-UW-2000; 2
05-UW-2000; 2
28-UUL-2000; 2
22-AUG-2000; 2
08-MOV-2000; 2
01-DEC-2000; 2
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PR 10-JUN-1999; 99US-0089722P.
PR 10-JUN-1999; 98US-0089732P.
PR 10-JUN-1999; 98US-008873P.
PR 10-JUN-1999; 98US-008882P.
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PR 23-JUN-1999; 98US-0099999; 98US-00999999.
PR 23-JUN-1999; 98US-00999999.
PR 23-JUN-199
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Gaps
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es 0; Indels
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17-OCT-1997;
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10.SEP-1998;
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01-OCT-1998,
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06-MAY-1998;
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tumour necrosis factor alpha; chondrocyte cell; tumour; gene therapy;
tissue typing.
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6; Conservative 0; Mismatches
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97US-005926P.
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97US-0063120P.
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97US-0063124P.
97US-0063544P.
97US-0063544P.
97US-0063544P.
      98WO-US019330.
98US-01006883P.
98US-0100919P.
98US-0100319P.
98US-0101014P.
98US-0101014P.
98US-01011471P.
98US-0101471P.
98US-0101473P.
98US-0101739P.
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Best Local Similarity
Matches 6; Conserv
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980S-0089538P.
980S-0089538P.
980S-0089563P.
980S-0089952P.
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Human; PRO; secreted protein; transmembrane protein; extracellular domain; tumour necrosis factor-alpha; TNF-alpha; chondrocyte; proliferation; differentiation; cartilage disorder; bone disorder; arthritis; sports injury; cancer; tumour; diagnosis; adrenal tumour; lung; colon; breast; prostate; kidney; rectum; cervix; liver; drug screening; transgenic animal; genetic analysis; antiarthritic; vulnerary; gene therapy.
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100.0%; Pred. No. 4.8
:ive 0; Mismatches
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980S-0102330P
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24-NOV-1997;
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Matches
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No. 4.8e+02;
O; Indels
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Pred. No.
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100.0%; Pred
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Best Local Similarity
Matches 6; Conserv
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21-0CT-1997;
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18-DEC-1997;
10-MAR-1998;
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tumour necrosis factor alpha; chondrocyte cell; tumour; gene therapy;
tissue typing.
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                                                                                                                                                                                                    0;
                                                                                                                                                        Similarity 100.0%; Score 6; DB 6; Length 1089; Similarity 100.0%; Pred. No. 4.8e+02; 6; Conservative 0; Mismatches 0; Indels
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98US-0102330P.
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98US-0102571P.
98US-0102684P.
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97US-0053266P
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                                                                                                                                                                             Local Similarity
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01-OCT-1998;
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17-OCT-1997

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08-APR-1998;
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                                                                                                                                                               Query Match
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Matches
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ABU99599
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24-JUN-1
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                                                0;
              Query Match
30.0%; Score 6; DB 6; Length 1089;
Best Local Similarity 100.0%; Pred. No. 4.8e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
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chromosome mapping; gene mapping; cytostatic.
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970S-0053466P.
970S-0063140P.
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                                                                                                                                                                                                                                                                                 Human PRO polypeptide #133.
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687 WLRRYG 692
                                                                             12 WLRRYG 17
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11-DEC-1997;
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18-DEC-1997;

10-MAR-1998;

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31-MAR-1998;

31-MAR-1998;

31-MAR-1998;

31-APR-1998;

31-APR-1998;
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ABU82838
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Human; gene therapy, tissue typing; tumour; chondrocyte proliferation; chondrocyte differentiation; tumour necrosis factor-alpha release; affinity purification.
                                                                                                                                                                                                  Novel human secreted and transmembrane protein PRO1249.
                                                                                          ABU89959 standard; protein; 1089 AA.
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9703-0063260P-
9703-0063486P-
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                                                                                                                                                                 (first entry)
     687 WLRRYG 692
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21-NOV-1997;
24-NOV-1997;
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12-DEC-1997;
17-DEC-1997;
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11-MAR-1998;
20-MAR-1998;
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06-MAY-1998;
07-MAY-1998;
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31-MAR-1998;
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98US-0103449P.
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98US-0103395P
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Best Local Similarity
Matches 6; Conserv
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02-JUL-1998;
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04-AUG-1998;
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01-SEP-1998;
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18-SEP-1998;
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23-SEP-1998;
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29-SEP-1998;
29-SEP-1998;
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01-OCT-1998;
01-OCT-1998;
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16-SEP-1998;
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17-SEP-1998;
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24-SEP-1998;
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PR 17-14X-1996 9815-00864640P.
PR 15-14X-1996 9815-00864640P.
PR 15-14X-1996 9815-008653PP.
PR 15-14X-1996 9815-008653PP.
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PR 22-14X-1996 9815-008623PP.
PR 22-14X-1996 9815-008623PP.
PR 02-14X-1996 9815-008622PP.
PR 10-14X-1996 9815-008622PP.
PR 10-14X-1996 9815-008622PP.
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PR 12-14X-1996 9815-009622PP.
PR 12-14X-1996 9815-009622PP.
PR 12-14X-1996 9815-009622PP.
PR 12-14X-1996 9815-009622PP.
PR 12-14X-1996 9815-00962PP.
PR 12-14X-1996 9815-009622PP.
PR 12-14X-1996 9815-00962PP.
PR 12-14X-1996 9815-00962PP.
PR 12-14X-1996 9815-00962PP.
PR 12-14
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Gaps
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Pred. No. 4.8
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 66
ABR68208
ID ABR68208 standard; protein; 1089 AA.
XX
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Similarity 100.0%; P
6; Conservative 0;
               98US-0096757P.
98US-0096757P.
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98US-0100684P.
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98US-010178BP.
 98US-0096012P
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687 WLRRYG
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10-SEP-1998;
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Matches
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9805-0085582P

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9805-0086392P

9805-008720BP

9805-008727P

9805-0087627P

9805-0087627P

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110-JUN-1998

110-JUN-1998
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02-JUL-1998;
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   extracellular domain, tumour necrosis factor alpha, TNP-alpha, chondrocyte, proliferation, differentiation, cartilage disorder; bone disorder, arthritis, sports injury, cancer, tumour, diagnosis, adrenal tumour, lung; colon, breast, prostate, kidney; rectum, cervix, liver, drug screening; transgenic animal; genetic analysis; antiarthritic, vulnerary; gene therapy.
                                                     Human; PRO; secreted protein; transmembrane protein;
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97US-0063121P.
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97US-0063541P.
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97US-0081334P.
97US-0081349P.
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97US-00844334P.
97US-00844334P.
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31-0CT-1297;
31-0CT-1297;
24-NOV-1997;
24-NOV-1297;
                                                                                                                             Homo sapiens.
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12-DEC-1997;
17-DEC-1997;
18-DEC-1997;
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29-APR-1998
                  11-AUG-2003
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ABR68208;
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knockout; chromosome identification; tissue typing; tumour; chondrocyte proliferation; chondrocyte differentiation; tumor necrosis factor-alpha release stimulator.
                                                                                                         970S-0059263P.
970S-0059266P.
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970S-0083149P.
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17-DEC-1997;
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07-MAY-1998;
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0
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Pred. No. 4.86
0; Mismatches
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100.0%; Fig.
98US-0096891P-
98US-0096931P-
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98US-0096939P-
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98US-0097952P-
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98US-010048P-
98US-010048P-
98US-010147P-
98US-0101739P-
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25-SEP-1998
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Matches
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98US-0098014P

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98US-0081025P

98US-0088028P

98US-0088033P

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98US-0088033P

98US-0088033P

98US-008812E

98US-008812E

98US-008812E

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Gaps
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hes 0;
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Conservative 0;
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es 6; Conserv
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Matches
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                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                  Query Match 30.0%; Score 6; DB 6; Length 1085
Best Local Similarity 100.0%; Pred. No. 4.8e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted/transmembrane protein (PRO) #133.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO08769 standard; protein; 1089 AA.
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Human; secreted and transmembrane protein; PRO; TNF-alpha; thumour necrosis factor alpha; chondrocyte cell; tumour; gene therapy; tissue typing; adrenal tumour; lung tumour; colon tumour; breast tumour; prostate tumour; rectal tumour; cervical tumour; liver tumour.
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nes 0; Indels
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Pred. No. 4.8e
0; Mismatches
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100.0%; Pre
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11-00N-1998;
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09-JUN-1998;
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No. 4.8e+02;
0; Indels
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Pred. No. 4.8e
0; Mismatches
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100.0%; Pre
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97US-0062250P.
97US-0063120P.
97US-0063121P.
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97US-0063541P.
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28-0CT-1997;
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29-0CT-1997;
31-0CT-1997;
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18-SEP-1997;
17-OCT-1997;
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980S-0089538P

980S-0089538P

980S-0089953P

980S-0089952P

980S-0090545P

980S-0090444P

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98US-0100681P.
98US-0100684P.
98US-0100930P.
98US-0100849P.
98US-010104P.
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26-AUG-1998;
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Pred. No. 4.8e+02;
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        98US-0101475P
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970S-0063541P.
970S-0063511P.
970S-0063734P.
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970S-0063734P.
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687 WLRRYG 692
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7US-00680179 8US-00774529 8US-00776329 8US-0077649P 8US-0079886P 8US-0079786P 8US-00801979 8US-00801979 8US-00801979 8US-00801979 8US-00801979 8US-00801979 8US-00801979	9805 0082568F 9805 0082568F 9805 0082568F 9805 0082769F 9805 0083229F 9805 0083429F 9805 0083459F 9805 0084640F 9805 0085709F 9805 008709F 9805 008709F 9805 0087058F 9805 0088025F 9805 0089025F 9805	805-00880399 805-00880399 805-00881679 805-0088205 805-00882179 805-00887125 805-00887259 805-00887389 805-00887389 805-00887409 805-00887409 805-00887409 805-00887409 805-00887409 805-00887409 805-00887409 805-00887409 805-00887409 805-00887409 805-00887618 805-00897389 805-00895389 805-00895389 805-00895389 805-00895389 805-00895389
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22-70N-1998,
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24-70N-1998,
25-70N-1998,
25-70

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Three hundred and five nucleic acids encoding PRO polypeptides, useful for the manufacture of a medicament for diagnosing or treating tumor.
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ith V, Watanabe CK, Wood WI,
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                                                                                                                                                          10-MAY-2001; 2001US-00854208.
25-MAY-2001; 2001US-00854208.
01-JUN-2001; 2001WS-0086028.
05-JUN-2001; 2001WS-00874503.
20-JUN-2001; 2001WS-00874503.
29-JUN-2001; 2001WS-US019692.
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18-JUL-2001; 2001US-00908827.

30-JUL-2001; 2001US-00918585.

06-AUG-2001; 2001US-00924419.

13-AUG-2001; 2001US-00931836.

28-AUG-2001; 2001US-00941992.

29-AUG-2001; 2001US-00941992.

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2000WO-US020710.
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2000US-00747259.
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N-PSDB; ACA72903.
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28-FEB-2001;
22-MAR-2001;
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18-SEP-2000;
08-NOV-2000;
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20-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                Human; PRO; secreted polypeptide; transmembrane polypeptide;
tumour necrosis factor alpha; TNF-alpha; chondrocyte cell; tumour;
                                                                                                     DB 6; Length 1089; 4.8e+02;
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                                                                                                                                                           687 WLRRYG 692
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Matches 6; Conserv
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01-DEC-1999;
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                                                                                                                                                                                                                                                                                                     cytostatic.
                                                                                                                                                                                                                           ABU85710;
                                                                                                                                                                                       RESULT 73
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Godowski PJ, Gurney AL;

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                        transmembrane polypeptides) and the PRO polypeptides 'setzeru and transmembrane polypeptides on a method for stimulating the release of The invention also relates to a method for stimulating the release of tumour necrosis factor alpha (TNF-alpha) from human blood by contecting the blood with a sequence of the invention, a method for stimulating the proliferation or differentiation of chondrocyte cells by contacting the proliferation or differentiation of chondrocyte cells by contacting the tumour in a mammal. The polypeptides and polymucleotides are useful for the manufacture of a medicament for diagnosing or treating a tumour in a mammal. Sequences ABUBSST8-ABUBSSB2 represent human PRO polypeptides of the invention. Note: The sequence data for this patent is also available in electronic format from USPTO at seqdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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The invention relates to human PRO polypeptides (secreted and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6; Length 1089;
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17-JUN-
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980S-008349SP
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ABU98870
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ACC ABU9
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Human; secreted and transmembrane protein; PRO; cytostatic; gene therapy; chondrocyte stimulator; tumour; adrenal tumour; lung tumour; colon tumour; breast tumour; prostate tumour; rectal tumour; cervical tumour; ilver tumour; chromosome identification.
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Pred. No. 4.8e+02;
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100.0%; Pre
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98US-0102687P.
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Matches 6
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Search completed: October 26, 2004, 07:16:10 Job time : 92 secs

us-10-066-965a-1.oligo.rapb

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6 US-10-437-963-199246

1 US-09-764-875-621

5 US-10-34-599-160000

5 US-10-38-12A-47856

4 US-10-389-493-4601

US-09-815-224-11699

US-09-815-247-11699

6 US-10-369-493-7358

6 US-10-369-161A-9

6 US-10-289-161A-9

6 US-10-289-161A-9

6 US-10-389-161A-9

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7 US-10-725-189-8

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10 US-10-725-189-8
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US-10-1052-586-266
US-10-1052-586-266
US-10-105-737-266
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US-10-105-737-266
US-10-105-737-266
US-10-105-737-266
US-10-105-737-266
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US-10-105-791-266
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Sequence 1, Appli
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Sequence 308, App
Sequence 1361.5,
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Sequence 1614, Ap
Sequence 194850,
Sequence 189867,
Sequence 183967,
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                                                                                  ; Search time 65 Seconds (without alignments) 99.618 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                               1364641
          GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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4 US-10-066-965A-1

4 US-10-066-965A-2

4 US-10-066-965A-2

4 US-10-066-965A-8

4 US-10-226-956-308

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5 US-10-424-599-252446

US-09-867-550-1614

US-10-424-599-183967

5 US-10-424-599-1833967

5 US-10-424-599-1833967

5 US-10-424-599-1833967
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                                                                                                                                                                                                                             1364641 seqs, 323758627 residues
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                                                                                                                                                                                                                                                                                                                                             Post-processing: Listing first 100 summaries
                                                          - protein search, using sw model
                                                                                     October 26, 2004, 07:22:37
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Gapop 60.0 , Gapext 60.0
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Word size :

Searched:

Sequence:

Database :

Sequence

11111 1211008765432

Result No. 0;

Gaps

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APPLICANT: COLAS, PIERRE
APPLICANT: REEMY, ROCBR
APPLICANT: REEMY, ROCBR
APPLICANT: COHEN, BARAK A.
TITLE OF INVENTION: TARGETED MODIFICATION OF INTRACELLULAR COMPOUNDS
FILE REPERENCE: BGYP 3.0-015
CURRENT APPLICATION NUMBER: US/10/066,965A
CURRENT FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: BRENT, ROGER
APPLICANT: COHEN, BARAK A.
TITLE OF INVENTION: TAGESTED MODIFICATION OF INTRACELLULAR COMPOUNDS
FILE REFERENCE: EGYP 3.0-015
CURRENT APPLICATION NUMBER: US/10/066,965A
CURRENT FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ) OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: peptide US-10-066-965A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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100.0%; Pred. No. 1.1e-13;
tive 0; Mismatches 0;
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100.0%; Pred. No. 1.1e-13;
tive 0; Mismatches 0;
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100.0%; Pred. No. 1.1e-13;
                Mismatches
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Publication No. US2003143626A1
GENERAL INFORMATION:
APPLICANT: COLAS, PIERRE
                                                                                                                                                                      Sequence 1, Application US/10066965A Publication No. US20030143626A1 GENERAL INFORMATION:
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                                                    OVWSLWALGWRWLRRYGWNM
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Best Local Similarity 100.
Matches 20, Conservative
              Conservative
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Matches 20; Conserv
 Best Local Similarity
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US-10-066-965A-7
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US-10-066-965A-1
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LENGIH: 20
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                                                      Sequence 266,
Sequence 266,
Sequence 266,
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                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xu, Chanxing Wilson TITLE OF INVENTION: INTERACTION TRAP SYSTEMS FOR DETECTING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 14; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: DESCRIPTION OF THE CONTRY: USA

COUNTRY: USA

CONPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/10/162,538

FILING DATE: 04-Jun-2002

CLASSIFICATION: CUNROWN>
PRIOR APPLICATION: CUNROWN>
PRIOR APPLICATION UMBER: US/08/630,052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: No. US20030113749A1 Relevant
US-10-176-487-266
US-10-176-493-266
US-10-176-913-266
US-10-176-919-266
US-10-176-919-266
US-10-176-925-266
US-10-176-925-266
US-10-180-543-266
US-10-180-543-266
US-10-180-543-266
US-10-180-546-266
US-10-180-547-266
US-10-180-547-266
US-10-180-547-266
US-10-180-547-266
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REFERENCE/DOCKET NUMBER: 00786/311001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                 ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
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APPLICATION NUMBER: 08/504,538
FILING DATE: July 20, 1995

APPLICATION NUMBER: 08/278,082
FILING DATE: July 20, 1994

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                    RESULT 102-538-12
US-10-162-538-12
Sequence 12, Application US/10162538
Publication No. US20030113749A1
GENERAL INFORMATION:
APPLICANT: Brent, Roger
GOSTON NO. USSON NO. USPAIRM H. USSON, TIMM H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INTERACTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 12: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 28 CORRESPONDENCE ADDRESS:
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   US-10-162-538-12
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APPLICANT: Brophy, Colleen
APPLICANT: Brophy, Colleen
APPLICANT: Formalavilas, Padmini
APPLICANT: Panitch, Alyssa
APPLICANT: Panitch, Alyssa
APPLICANT: Joshi, Lokesh
APPLICANT: Joshi, Lokesh
APPLICANT: Joshi, Lokesh
APPLICANT: Saal, Brandon L.
TITLE OF INVENTION: REAGENTS AND METHODS FOR SMOOTH MUSCLE THERAPIES
FILE REFERENCE: ASU-1061-08
CURRENT APPLICATION NUMBER: 105/10/226,956
CURRENT FILING DATE: 2002-08-23
PRIOR APPLICATION NUMBER: 60/314,535
PRIOR FILING DATE: 2001-08-23
NUMBER OF SEQ ID NOS: 320
SUSTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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TITLE OF INVENTION: REAGENTS AND METHODS FOR SMOOTH MUSCLE THERAPIES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 14;
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100.0%; Pred. No. 28;
Live 0; Mismatches
                      FILE REFERENCE: ASU-1061-US
CURRENT PEDFLICATION NUMBER: US/10/226,956
CURRENT FILING DATE: 2002-08-23
PRIOR APPLICATION NUMBER: 60/314,535
PRIOR FILING DATE: 2001-08-23
NUMBER OF SEQ ID NOS: 320
SOFTWARE: PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-437-963-136125
; Sequence 136125, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; LOCATION: (20)...(20); CTHER INFORMATION: S is phosphorylated US-10-226-956-305
                                                                                                                                                                                                                                                                                                                                                                                       ; LOCATION: (17)...(17); OTHER INFORMATION: S is phosphorylated US-10-226-956-308
                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Synthetic peptide
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ORGANISM: Artificial sequence
FBATURE:
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Best Local Similarity
Matches 6; Conserv
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Publication No. US20030143626A1
GENERAL INFORMATION:
BERRE
APPLICANT: COLAS, PIERRE
APPLICANT: COLAS, BERRE
APPLICANT: RRENT, ROGER
APPLICANT: TARGETED MODIFICATION OF INTRACELLULAR COMPOUNDS
FILE REPERENCE: EGYP 3.0-0.15
CURRENT APPLICATION NUMBER: US/10/066,965A
CURRENT FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Patentin Ver. 2.1
                                                                                        APPLICANT: COLAS, PIERRE
APPLICANT: COLAS, PIERRE
APPLICANT: COGEN
APPLICANT: COHEN, BARAK A.
TITLE OF INVENTION: TARGETED MODIFICATION OF INTRACELLULAR COMPOUNDS
FILE REFERENCE: EGYP 3.0-0.15
CURRENT APPLICATION NUMBER: US/10/066,965A
CURRENT FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       65.0%; Score 13; DB 14; I 100.0%; Pred. No. 1.6e-06;
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Best Local Similarity 100.0%; Pred. No. 1.6e-06;
Matches 13; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 308, Application US/10226956
Publication No. US20030060399A1
GENERAL INFORMATION:
APPLICANT: Brophy, Colleen
APPLICANT: Komalavilas, Padmini
APPLICANT: Panitch, Alyssa
APPLICANT: Joshi, Lokesh
APPLICANT: Seal, Brandon L.
                    Sequence 2, Application US/10066965A Publication No. US20030143626A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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US-10-066-965A-8
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LENGTH: 20
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LENGTH: 20
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Matches

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TITLE OF INVENTION: No. US20120081206Alel Polymucleotides from Atherogenic Cells and TITLE OF INVENTION: Thereby FILE REFERENCE: 21402-013 (Cura-313)
FILE REFERENCE: 1402-013 (Cura-313)
CURRENT APPLICATION NUMBER: US50/967,550
CURRENT FILING DATE: 2001-09-20
PRIOR APPLICATION NUMBER: USSN 60/208,427
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 2125
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Li, Ping,
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5321)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 194850
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US-10-437-963-194850
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100.0%; Pred. No. 85;
iive 0; Mismatches
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LOCATION: (1)..(120)
OTHER INFORMATION: unsure at all Xaa locations
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100.0%; Pred. No. 81;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 194850, Application US/10437963; Publication No. US20040123343A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | Sequence 183967, Application US/10424599 | Publication No. US20040031072A1 | GENERAL INFORMATION: APPLICANT: La Rosa Thomas J | APPLICANT: Kovalic David K | APPLICANT: Zhou Yihua | APPLICANT: Cao Yongwei | APPLICANT: Ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
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Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 6; Conserv
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US-10-437-963-194850
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US-10-424-599-183967
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                                                            APPLICANT: Zhou, Yihua APPLICANT: Zhou, Yihua APPLICANT: Zhou, Yihua APPLICANT: Cao, Yongwei APPLICANT: Wu, Wei Hei APPLICANT: Burkarov, Andrey A. APPLICANT: Burkarov, Andrey A. APPLICANT: Barbazuk, Brad APPLICANT: Li, Ping APPLICANT: Li, Ping TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 136125
LENGTH: 65
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Publication No. US20040031072A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPRESENCE: 38-21(5323)8
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
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US-10-424-599-252446
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US-10-437-963-136125
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Fatent No. US20206A1
GENERAL INFORMATION:
APPLICANT: Leach, Martin D.
APPLICANT: Mehraban, Fuad,
APPLICANT: Conley, Pamela
APPLICANT: Law, Debbie
APPLICANT: Topper, James
   La Rosa, Thomas J.
Kovalic, David K.
Zhou, Yihua
Cao, Yongwei
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ORGANISM: Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10 WRWLRR 15
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US-10-424-599-252446
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US-09-867-550-1614
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LENGTH: 74
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Gaps

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APPLICANT: Li, Fing TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Rice Nucleic Acid Molecules and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B CURRENT APPLICATION NUMBER: US/10/437,963 CURRENT FILING DATE: 2003-05-14 NUMBER OF SEQ ID NOS: 204966
  TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFRENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73129
SEQ ID NO 40080
LENGTH: 196
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US-10-437-963-199236
                                                                                                                                                                                                                                                                                                              30.0%; Score 6; DB 15; Le 100.0%; Pred. No. 1.2e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: unsure
LOCATION: (1)..(202)
OTHER INFORMATION: unsure at all Xaa locations
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US-10-425-114-40080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 199236, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
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Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Avoralic, David K.
APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Boukharov, Andrey A.
                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity luv...
5. Conservative
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ORGANISM: Oryza sativa
                                                                                                                                                                     TYPE: PRT
ORGANISM: Glycine max
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US-10-437-963-199236
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LENGTH: 202
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APPLICANT:
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APPLICANT:
APPLICANT:
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Fublication No. US20040034898A1

GENERAL INFORMATION:

APPLICANT: Liu, Jingdong

APPLICANT: Screen, Steven B

APPLICANT: Screen, Steven B

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

TILE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

TILE OF UNENT APPLICATION NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

SEQ ID NO 43111

LENGTH: 180
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With FILLS OF INVENTION: Plants and Uses Thereof for Plant Improvement FILLS REFERENCE: 38-21(53223)B CURRENT APPLICATION NUMBER: US/10/424,599 CURRENT FILING DATE: 2003-04-28 NUMBER OF SEQ ID NOS: 285684

ERROR THE SEQ ID NOS: 285684

LENGTH: 180
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
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Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Clone ID: 700735202_FLI.pep
US-10-425-114-43111
                                                                                                                                                                                                                                                                                        LOCATION: (1)..(180)
OTHER INFORMATION: unsure at all Xaa locations
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100.0%; Pred. No. ...
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US-10-425-114-40080
Sequence 40080, Application US/10425114
Sequence 40080, Application US/10425114
Sequence 10080, Application US/10425114
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
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APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao Vormeri
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Best Local Similarity 100.
Matches 6; Conservative
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                                                                                                                                                                                                TYPE: PRT ORGANISM: Glycine max
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Matches 6; Conserva
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LOCATION: (1)..(1
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Gaps

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LOCATION: (29)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
NAME/KEY: SITE
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INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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CURRENT APPLICATION NUMBER: US/09/764,875
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1249
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 621
LENGTH: 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30.0%; Score 6; DB 11; Length 305; 100.0%; Pred. No. 1.7e+02; ive 0; Mismatches 0; Indels
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Sequence 4, Application US/10343650A

Sequence 4, Application US/10343650A

Publicaturin No. US20040067499A1

GENERAL INFORMATION:

APPLICANT: HAGA, TATSUYA

TITLE OF INVENTION: NOVEL G-PROTEIN COUPLED RECEPTOR

FILE REFERENCE: 31671-186347

CURRENT APPLICATION NUMBER: US/10/343,650A

CURRENT FILING DATE: 2003-07-21

PRIOR APPLICATION NUMBER: JP 2000/237818

PRIOR APPLICATION NUMBER: JP 2000/34434

PRIOR FILING DATE: 2001-02-13

NUMBER OF SEQ ID NOS: 694

SOFTWARE: Patentin Ver. 2.1
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 6; Conserv
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Best Local Similarity
Matches 6; Conserv
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US-10-424-599-160000
                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: SITE
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OTHER INFO
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; Publication No. U520040123343A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Cao, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Bribazuk, Brad
APPLICANT: Bribazuk, Brad
APPLICANT: Bribazuk, Brad
APPLICANT: Bribazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT APPLICATION NUMBER: US/10/437,963
SRQ ID NO 199246
; SRQ ID NO 199246
                   APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and USes Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53221) B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 134324
LENGTH: 240
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Sequence 621, Application US/09764875

Publication No. US20040018969A1

GENERAL INFORMATION:

APPLICANT: Rosen et al.

TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies

FILE REFERENCE: PJZ02
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US-10-437-963-134324
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US-10-437-963-199246
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LOCATION: (1)..(243)
OTHER INFORMATION: unsure at all Xaa locations
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Best Local Similarity 100...
Thes 6; Conservative
    Barbazuk, Brad
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                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Oryza sativa
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Best Local Similarity
Matches 6; Conserv
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US-10-437-963-199246
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Sequence 4601, Application US/10369493
Publication No. US2003023367SA1
GENERAL INFORMATION:
APPLICANT: Cao, Yongwei
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
FILE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052) B
CURRENT APPLICATION NUMBER: US/10/369,493
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PRIOR FILING DATE: 2001-02-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 78614
SOFTWARE: Patentin version 3.1
SEQ ID NO 47856
LENGTH: 396
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100.0%; Pred. No. 2.1e+02;
cive 0; Mismatches 0;
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APPLICANT: HORIKAWA, JUN
APPLICANT: SHIBA, TADANOSHI
APPLICANT: SHIBA, TADANOSHI
APPLICANT: SHIBA, TADANOSHI
APPLICANT: HATTORI, WOSHIRA
TITLE CANT: HATTORI, MASAHIRA
TITLE REFERENCE: 249-262
CURRENT APPLICATION NUMBER: US/10/156,761
CURRENT PILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: JP 2001-204089
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR FILING DATE: 2001-08-02
NUMBER: OF SEQ ID NOS: 15109
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PRIOR FILING DATE: 2002-02-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-156-761-12394
; Sequence 12394, Application US/10156761
; Publication No. US2030119018A1
GENERAL INFORMATION:
; APPLICANT: OWURA, SATOSHI
; APPLICANT: IKEDA, HARUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Streptomyces avermitilis
                                                                                                                                                                     TYPE: PRT
ORGANISM: Burkholderia cepacia
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Best Local Similarity 100.
Matches 6; Conservative
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US-10-369-493-4601
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                                             APPLICANT: La Rosa Thomas J
APPLICANT: La Rosa Thomas J
APPLICANT: La Rosa Thomas J
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: 18-21 (53223) B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT APPLICATION DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 160000
LENGTH: 369
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Pred. No. 2e+02;
0; Mismatches 0; Indels
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US-10-424-599-160000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: unsure
LOCATION: (1)..(369)
OTHER INFORMATION: unsure at all Xaa locations
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CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
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Sequence 47856, Application US/10282122A
Publication No. US20040029129A1
SEQUENCE APPLICANT: APPLICANT: APPLICANT: Malone, Carlos
APPLICANT: Malone, Carlos
APPLICANT: Haselbeck, Robert
APPLICANT: Haselbeck, Robert
APPLICANT: Wall, Daniel
APPLICANT: Wall, Daniel
APPLICANT: Yamanoto, ApplicANT: APPLICANT: Yamanoto, ApplicANT: Trawick, John
APPLICANT: Yamanoto, Robert
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PRIOR APPLICATION NUMBER: 60/191,078
PRIOR PILING DATE: 2000-02-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR PELING DATE: 2000-05-23
PRIOR PELING DATE: 2000-05-23
PRIOR PELING DATE: 2000-05-26
PRIOR PELING DATE: 2000-09-6
PRIOR PELING DATE: 2000-09-6
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR APPLICATION NUMBER: 60/230,347
PRIOR PELING DATE: 2000-9-6
PRIOR PELING DATE: 2000-9-6
PRIOR PELING DATE: 2000-10-23
PRIOR PELING DATE: 2000-10-23
PRIOR PELING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/25,362
PRIOR APPLICATION NUMBER: 60/25,362
PRIOR APPLICATION NUMBER: 60/25,361
PRIOR APPLICATION NUMBER: 60/25,391
PRIOR PELING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/25,391
PRIOR PELING DATE: 2000-12-22
PRIOR PELING DATE: 2000-12-29
PRIOR PELING DATE: 2000-12-29
PRIOR PELING DATE: 2001-02-09
PRIOR PELING DATE: 2001-02-09
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100.0%;
  Publication No. US20040031072A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0
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ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 268 LRRYGW 273
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Gaps

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APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goodman, Barry S.
APPLICANT: Goodman, Barry S.
APPLICANT: Gregory J.
APPLICANT: Chark WITH IMPROVED PROPERTIES
FILE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REPERRECE: 38-10/52052)
FILE REPERRECE: 38-10/52052)
FILE REPERRECE: 2003-02-28
PRIOR FILING DATE: 2003-02-28
PRIOR FILING DATE: 2002-02-21
NUMBER OF SRQ ID NOS: 47374
SRQ ID NO 7358
LENGTH: 429
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No. 2.2e+02;
0; Indels
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LOCATION: (1)..(429)
OTHER INFORMATION: unsure at all Xaa locations
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100.0%; Pred. No. 2.2
tive 0; Mismatches
                     CURRENT APPLICATION NUMBER: US/10/389,647
CURRENT FILING DATE: 2003-03-14
PRIOR APPLICATION NUMBER: 09/653730
PRIOR FILING DATE: 2000-09-01
PRIOR PLING DATE: 1999-09-03
NUMBER OF SEQ ID NOS: 710
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 7358, Application US/10369493; Publication No. US20030233675A1; GENERAL INFORMATION:
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
APPLICANT: Ww, Wei
APPLICANT: Ww, Wei
APPLICANT: Barbazuk, Brad
                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Burkholderia cepacia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Conservative
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Matches 6; Conservative
FILE REFERENCE: UIZ-038CP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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US-10-437-963-122938
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US-10-369-493-7358
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                                                                                                                                                                             Length 400
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                                                                                                                                                                                                                                    0; Indels
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APPLICANT: Yamamoto, Robert T.
APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 575, Application US/10389647
CENDICALION NO. US20040033549A1
CENERAL INFORMATION:
APPLICANT: GREENBERG, E. Peter
APPLICANT: SCHUCKFR, Martin
APPLICANT: LOSTROH, Candi
TITLE OF INVENTION: QUORUM SENSING SIGNALING IN BACTERIA
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                                                                                                                                                                    30.0%; Score 6; DB 1
100.0%; Pred. No. 2.1
:ive 0; Mismatches
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CURRENT FILING DATE:

PRIOR PILING DATE:

PRIOR PILING DATE:

PRIOR PILING DATE:

PRIOR FILING DATE:

PRIOR FILING DATE:

PRIOR FILING DATE:

PRIOR PILING DATE:

PRIOR PAPLICATION NUMBER: 60/206,848

PRIOR PILING DATE:

PRIOR PILING DATE:

PRIOR PILING DATE:

PRIOR PILING DATE:

PRIOR FILING DATE:

PRIOR PILING DATE:

PRI
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                                                   TYPE: PRT
ORGANISM: Burkholderia fungorum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Klebsiella pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ohlsen, Kari L.
Zyskind, Judith W.
Wall, Daniel
Trawick, John D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
                                                                                                                                                                    Query Match 30.0
Best Local Similarity 100.
Matches 6; Conservative
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Matches 6, Conserv
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                                                                                           RESULT 25
US-10-389-647-575
SEQ ID NO 4601
                              LENGTH: 400
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APPLICANT:
APPLICANT:
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Gaps

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APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221) B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 122935
LENGTH: 500
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APPLICANT: Turano, Frank J.
TTLE OF INVENTION: Methods for Regulating Plant GABA Production
FITLE OF INVENTION: Methods for Regulating Plant GABA Production
FILE REFERENCE: 722.45
CURRENT APPLICATION NUMBER: US 60/246,367
PRIOR APPLICATION NUMBER: US 60/246,367
PRIOR FILING DAIE: 2000-11.07
NUMBER OF SEQ ID NOS: 24
SSCTUMER: Patentin version 3.1
SEQ ID NO 2
LENGTH: 502
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APPLICANT: WALTERS, DANA M
APPLICANT: WALTERS, DANA M
APPLICANT: RAINER, RUSS
TITLE OF INVENTION: Genes Encoding Picric Acid Degradation
FILE REPERENCE: BC10.22 US NA
CURRENT APPLICATION NUMBER: US/09/955,597
CURRENT FILING DATE: 2001-09-17
                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Clone ID: PAT_MRT4530_25818C.1.pep US-10-437-963-122935
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100.0%; Pred. No. 2.4e+02;
iive 0; Mismatches 0;
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100.0%; Pred. No.
ative 0; Mismatch
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PRIOR FILING DATE: 1999-10-03
NUMBER OF SEQ ID NOS: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , Sequence 2, Application US/10006852; Publication No. US20030046732A1; GENERAL INFORMATION:
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; Sequence 3, Application US/09955597
; Publication No. US20020042117A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ; ORGANISM: Arabidopsis thaliana US-10-006-852-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Microsoft Office 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.08
Matches 6; Conservative
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US-10-006-852-2
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LENGTH: 532
                                                                                                                                                                                                                                                                 TYPE: PRT
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APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)8
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT APPLICATION NUMBER: US/10/437,963
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 122338
LENGTH: 472
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Sublication No. US20040023343A1

Sublicant: Balax: Richard H

APPLICANT: Broughton, Mary C

APPLICANT: Treadway, Patti J

APPLICANT: Treadway, Patti J

APPLICANT: Treadway, Patti J

APPLICANT: Treadway, Patti J

APPLICANT: Wadduri, Krishnamurthy

FILE REFERENCE: 50489 DIVI

CURRENT APPLICATION NUMBER: BALLER APPLICATION NUMBER: US/09/603,2078

PRIOR FILING DATE: EARLIER FILING DATE: 1998-03-09

SOFTWARE: PATENTING DATE: EARLIER FILING DATE: 1998-03-09

SOFTWARE: PATENTING DATE: PATENTER FILING DATE: 1998-03-09

SOFTWARE: PATENTING DATE: PATENTER FILING DATE: 1998-03-09

SOFTWARE: PATENTER FILING DATE: PATENTER FILING DATE: 1998-03-09

SOFTWARE: PATENTER FILING DATE: PATENTER FILING DATE: 1998-03-09

SOFTWARE: PATENTER FILING DATE: PATENTER FILING DATE: LIGHTHY

LIGHTHY 486
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No. 2.4e+02;
0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                              30.0%; Score 6; DB 16; Length 472; 100.0%; Pred. No. 2.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                       ), OTHER INFORMATION: Clone ID: PAT_MRT4530_25820C.1.pep
US-10-437-963-122938
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100.0%; Pred. No. 2.4
tive 0; Mismatches
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Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Rovalic, David K.
APPLICANT: Cao, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT; ORGANISM: Saccharopolyspora spinosa US-10-329-148A-16
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Best Local Similarity 10v...
5. Conservative
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                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         369 LRRYGW 374
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Matches 6; Conserv
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US-10-329-148A-16
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LENGTH: 548
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US-10-725-189-8
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   RESULT 34
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Eublication No. US20030152970A1
GENERAL INFORMATION:
APPLICANT: Millennium Pharmaceuticals, Inc.
APPLICANT: Silos-Santiago, Inmacculada
ITILE OF INVENTION: Methods and Compositions to Treat Pain
TITLE OF INVENTION: and Painful Disorders Using 577, 20739 or 57145
FILE REPERENCE: MPI2001-287P1R(M)
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                                                   30.0%; Score 6; DB 9; Length 532; 100.0%; Pred. No. 2.6e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/10308163
Publication No US20040106147A1
GENERAL INFORMATION:
APPLICANT: UCB, S.A.
APPLICANT: INNCE, Berkley
APPLICANT: FUCE, Rand
TITLE OF INVENTION: Levetiracetam Binding Site
TITLE OF INVENTION: Levetiracetam Binding Site
CURRENT APPLICATION NUMBER: US/10/308,163
CURRENT APPLICATION NUMBER: US/10/308,163
SOFTWARE: Patentin Version 3.1
SEQ ID NO 8
ORGANISM: Rhodococcus erythropolis HL PM-1
                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/289,161A
CURRENT FILING DATE: 2003-03-25
PRIOR APPLICATION NUMBER: US 60/333,073
PRIOR FILING DATE: 2001-11-06
NUMBER OF SEQ ID NOS: 9
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO
                                                 Query Match
Best Local Similarity 100.
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Best Local Similarity 100.
Matches 6; Conservative
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Matches 6; Conservative
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ORGANISM: Homo Sapiens
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US-10-308-163-8
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US-10-289-161A-9
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US-10-308-163-8
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US-09-955-597-3
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Sequence 8, Application US/10725189

Sequence 8, Application US/10725189

Fublication No. US20040204388A1

GENERAL INFORMATION:

APPLICANT: UCNS, S.A.

APPLICANT: LXNCH, Berkley

APPLICANT: LXNCH, Berkley

TITLE OF INVENTION: Beilzures, neurological diseases, endocrinopathies and hormonal

TITLE OF INVENTION: asizures, neurological diseases, endocrinopathies and hormonal

TITLE OF INVENTION: UNMBER: US/10/725,189

CURRENT APPLICATION NUMBER: US/20/32,129

PRIOR APPLICATION NUMBER: US 60/430,372

PRIOR APPLICATION NUMBER: US 60/506,764

PRIOR FILING DATE: 2003-12-02

PRIOR FILING DATE: 2003-19-30
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US-10-308-163-16

Sequence 16, Application US/10308163

Publication No. US20040106147A1

GENERAL INFORMATION:

APPLICANT: UCB, S.A.

APPLICANT: LYNCH, Berkley

APPLICANT: FUKS, Brun

TITLE OF INVENTION: Levetiracetam Binding Site

FILE REPRENCE: S3259-5007

CURRENT APPLICATION NUMBER: US/10/308,163
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Pred. No.
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US-10-725-189-16
Sequence 16, Application US/10725189
Publication No. US20040204388A1
GENERAL INFORMATION:
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100.0%; Pre
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 16
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SEQ ID NO 8
LENGTH: 548
                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Rattus norvegicus US-10-308-163-16
                                                                                                                                                                                                                                                                                                                                                                                                                                           6; Conservative
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APPLICANT: LYNCH, Berkley
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US-10-725-189-8
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Buckharov, Andrey A.
APPLICANT: Buckharov, Andrey A.
APPLICANT: Buckharov, Andrey A.
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53221) B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 196644
LENGTH: 666
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US-10-437-963-117634
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US-10-437-963-196644
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Pred. No. 2.9e+02;
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OTHER INFORMATION: unsure at all Xaa locations
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OTHER INFORMATION: unsure at all Xaa locations
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                                                      Sequence 117634, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
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Best Local Similarity 100...
6; Conservative
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Best Local Similarity
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NAME/KEY: unsure
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                              US-10-437-963-117634
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APPLICANT: Milenmium Pharmaceuticals, Inc.
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Baisof, Scott D.
APPLICANT: Eniasof, Scott D.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING
TITLE OF INVENTION: Hall APPLICANT: Bilasof, 53043, 21425, 27410, 38554, 38555, 55063, 57145, 59914,
TITLE OF INVENTION: HALNO PAINFUL DISORDERS USING 16386,
TITLE OF INVENTION: 194921, 18652, 33260, 58573, 30911, 85913, 14303, 16816,
TITLE OF INVENTION: 194921, 18652, 33260, 58573, 30911, 85913, 14303, 16816,
TITLE OF INVENTION: 10427 OR 32620
FILE REFERENCE: MOUNERR: US 60/444,781
FRICE REPLIANG DATE: 2003-03-04
FRICE PLING DATE: 2003-04-16
FRICE PLING DATE: 2003-05-04
FRICE PLING DATE: 2003-05-04
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FRICE PRICE FABLE FABLESQ FOR WINDERR: US 60/491,048
FRICE PLING DATE: 2003-07-30
FRICE PRICE FABLESQ FOR WINDOWS WE SECOND TO NOT A S
             AFPLICANT: FUKS, Bruno
TITLE OF INVENTION: Methods for the identification of agents for the treatment of
TITLE OF INVENTION: Seizures, neurological diseases, endocrinopathies and hormonal
TITLE OF INVENTION: diseases
FILE REFERENCE: 53529-5007-01-US
CURRENT APPLICATION NUMBER: US/10/725,189
CURRENT FILING DATE: 2003-12-02
PRIOR APPLICATION NUMBER: US 60/430,372
PRIOR FILING DATE: 2002-12-03
PRIOR FILING DATE: 2000-09-30
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1
SEQ ID NO 16
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Publication No. US20040204359A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Rattus norvegicus
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Best Local Similarity 100.
Matches 6; Conservative
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US-10-768-158-24
APPLICANT: NOCKA, Karl
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LENGTH: 548
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic ITTLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2830P1C1
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                                                                                                                                                                               Length 1083;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/946,374 CURRENT FILING DATE: 2001-09-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR PILLING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098713
PRIOR PILLING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098749
PRIOR PILLING DATE: 1998-09-01
PRIOR PELING DATE: 1998-09-01
PRIOR PELING DATE: 1998-09-02
PRIOR PILLING DATE: 1998-09-02
PRIOR PILLING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/098813
PRIOR PILLING DATE: 1998-09-02
PRIOR PILLING DATE: 1998-09-02
PRIOR PILLING DATE: 1998-09-02
PRIOR PILLING DATE: 1998-09-02
PRIOR PILLING DATE: 1998-09-09
PRIOR FILLING DATE: 1998-09-09
PRIOR PILLING DATE: 1998-09-09
                                                                                                    ; TYPE: PRT
; ORGANISM: Corynebacterium glutamicum
US-09-738-626-6914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ferrara, Napoleone
Fong, Sherman
Gao, Wei-Qiang
Goddard, Audrey
Godowski, Paul J.
Grimaldi, Christopher J.
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. Sequence 102, Application US/09946374
. Publication No. US20030073129A1
. GENERAL INFORMATION:
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APPLICATION NUMBER: 60/099642
FILING DATE: 1998-09-09
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Tumas, Daniel
Watanabe, Colin K.
Williams, P. Mickey
Wood, William I.
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Paoni, Nicholas F.
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Hillan, Kenneth J.
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Smith, Victoria
  PRIOR FILING DATE: 2000-08-03
                  NUMBER OF SEQ ID NOS: 7059
SOFTWARE: PatentIn ver. 3.0
SEQ ID NO 6914
LENGTH: 1083
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Eaton, Dan L.
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Best Local Similarity
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  0; Indels
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100.0%; Pred. No. ...
  0; Mismatches
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PRIOR PLING DATE: 2000-12-18

PRIOR FILING DATE: 1999-12-16

PRIOR PLILING DATE: 1999-12-16

PRIOR APPLICATION NUMBER: UP 00/159162

PRIOR FILING DATE: 2000-04-07

PRIOR PLILING DATE: 2000-08-03

NUMBER OF PRIOR DATE: 2000-08-03

NUMBER OF PRIOR DATE: 2000-08-03

NUMBER OF PROBATE: 2000-08-03

SOFTWARE: PARENTIN VOR: 3.0

SEQ ID NO 3935

LENGTH: 755
                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES FILE REFERENCE: 249-125 CURRENT APPLICATION NUMBER: US/09/738,626
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APPLICANT: 0ZAKI, AKTO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
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PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 1999-12-16
PRIOR PPLICATION NUMBER: JP 00/159162
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: JP 00/280988
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; Sequence 6914, Application US/09738626
; Publication No. US20020197605A1
; GENEAL INFORMATION:
; APPLICANT: NAKAGAWA, SATOSHI
                                                                                                                                                                            Sequence 3935, Application US/09738626; Publication No. US20020197605A1; Publicant INFORMATION: APPLICANT: NAKAGAWA, SATOSHI
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ANDO, SEIKO
HAYASHI, MIKIRO
OCHIAI, KEIKO
YOKOI, HARUHIKO
TATEISHI, NAOKO
SENOH, AKIHIRO
IKEDA, MASATO
OZAKI, AKIO
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ANDO, SEIKO
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CCHIAL, KEKO
YOKOI, HAWUHIKO
TATEISHI, NAOKO
SENOH, AKTHIRO
IKEDA, MASATO
OZAKI, AKI
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Best Local Similarity 100.
Matches 6; Conservative
  6; Conservative
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NUMBER: 60/099741 1998-09-10 NUMBER: 60/099754 1998-09-10 NUMBER: 60/099763 1998-09-10 NUMBER: 60/099792 1998-09-10 NUMBER: 60/099808 1998-09-10 NUMBER: 60/099812 1998-09-10 NUMBER: 60/099812 1998-09-10 NUMBER: 60/099815	NUMBER: 60/100385 1999-09-15 1998-09-15 NUMBER: 60/100388 11998-09-15 NUMBER: 60/100584 11998-09-16 NUMBER: 60/10067 11998-09-16 NUMBER: 60/100661 11998-09-16 NUMBER: 60/100661 11998-09-16 NUMBER: 60/100664 11998-09-16 NUMBER: 60/100683 11998-09-17 NUMBER: 60/100684 11998-09-17 NUMBER: 60/100684 11998-09-17 NUMBER: 60/100684 11998-09-17 NUMBER: 60/100710 11998-09-17 NUMBER: 60/100710 11998-09-17 NUMBER: 60/100710	11998-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-1 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2 10988-09-2	1998-09- NUMBER: 6 NUMBER: 6 NUMBER: 6 1998-09- NUMBER: 6
APPLICATION DATE: APPLICATION DATE: APPLICATION DATE: FILING DATE: APPLICATION FILING DATE: APPLICATION FILING DATE: APPLICATION PILING DATE: APPLICATION FILING DATE: APPLICATION DATE: APPLICATION DATE: APPLICATION FILING DATE: APPLICATION DATE: APPLICAT	APPLICATION PILLING DATE: APPLICATION FILLING DATE: APPLICATION FILLIN	PILING DATE: APPLICATION N FILING DATE: APPLICAT	FILING DATE: APPLICATION FILING DATE: PELING DATE: APPLICATION FILING DATE: APPLICATION FILING DATE: APPLICATION
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PRIOR PLING DATE: 1998-09-24
PRIOR APPLICATION WURBER: 60/101915
PRIOR APPLICATION NUMBER: 60/101916
PRIOR APPLICATION NUMBER: 60/101916
PRIOR PLILING DATE: 1998-09-24
PRIOR PLILING DATE: 1998-09-29
PRIOR PLILING DATE: 1998-09-30
PRIOR PLILING DATE: 1998-10-07
PRIOR PLILING DATE: 1998-10-02
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PRIOR FILING DATE: 1997-12-18

PRIOR APPLICATION NUMBER: 60/077630

PRIOR APPLICATION NUMBER: 60/077632

PRIOR APPLICATION NUMBER: 60/077649

PRIOR FILING DATE: 1998-03-11

PRIOR FILING DATE: 1998-03-20

PRIOR FILING DATE: 1998-03-20

PRIOR PELING DATE: 1998-03-27

PRIOR PELING DATE: 1998-04-01

PRIOR PELING DATE: 1998-04-02

PRIOR PELING DATE: 1998-04-03

PRIOR PELING DATE: 1998-04-22

PRIOR PELING DATE: 1998-04-29

PRIOR PELING DATE: 1998-04-29
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APPLICATION NUMBER: 60/083559
FILING DATE: 1998-04-29
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PRIOR APPLICATION NUMBER: 60/084366
PRIOR FILING DATE: 1998-05-05
PRIOR PELLICATION NUMBER: 60/084414
PRIOR PLING DATE: 1998-05-06
PRIOR APPLICATION NUMBER: 60/084639
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PRIOR APPLICATION WUMBER: 60/084640
PRIOR FILING DATE: 1998-05-07
PRIOR APPLICATION NUMBER: 60/084643
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APPLICATION NUMBER: 60/086392
FILING DATE: 1998-05-22
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APPLICATION NUMBER: 60/085573
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FILING DATE: 1998-05-15
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FILING DATE: 1998-05-15
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APPLICANT: Wood, William I.

TITLE OF INVENTION: SCREETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: SCREETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: SCREETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC CURRENT PELICATION NUMBER: 05/05263

PRIOR PELICATION NUMBER: 05/05266

PRIOR PLING DATE: 1997-01-15

PRIOR PLING DATE: 1997-01-16

PRIOR PLING DATE: 1997-10-24

PRIOR PLING DATE: 1997-10-24

PRIOR PLING DATE: 1997-10-24

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PRIOR PLING DATE: 1997-11-24

PRIOR PLING DATE: 19
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                                                                       30.0%; Score 6; DB 10; Length 1089; 100.0%; Pred. No. 4.3e+02; ive 0; Mismatches 0; Indels
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Publication No. US20020127584A1
GENERAL INFORMATION:
   PRIOR APPLICATION NUMBER: 60/105807
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APPLICATION NUMBER: 60/069870
FILING DATE: 1997-12-17
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
Zhang, Zemin
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin I.
                                   Query Match
Best Local Similarity 100..
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430RTC1014
CURRENT APPLICATION NUMBER: US/10/176,758
CURRENT FILING DATE: 2002-06-21
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: SECRETED AND THE SAME
FILE REFERENCE: P3430R1642
CURRENT APPLICATION NUMBER: US,10/174,590
CURRENT APPLICATION NUMBER: US,10/174,590
CURRENT APPLICATION NUMBER: OS 2002-06-18
Prior application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LEDITH: 1089
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                                                                                        US-10-174-590-266
; Sequence 266, Application US/10174590
; Publication No. US20030008352A1
; GENERAL INFORMATION:
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Godowski, Paul J.
Gurney, Austin L.
Pan, James
Smith, Victoria
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Watanabe, Colin K,
Wood, William I.
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Best Local Similarity 100.
Matches 6; Conservative
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Goddard, Audrey
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CORGANISM: Homo Sapien
US-10-176-758-266
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US-10-174-590-266
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      687 WLRRYG 692
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100.0%; Pred. No. 4.3
tive 0; Mismatches
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FILING DATE: 1998-06-11
APPLICATION NUMBER: 60/08876
FILING DATE: 1998-06-11
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PRIOR FILING DATE: 1998-06-16
PRIOR PPLICATION NUMBER: 60/089538
PRIOR FILING DATE: 1998-06-17
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FILING DATE: 1998-06-17
LING DATE: 1998-05-22
PPLICATION NUMBER: 60/087098
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FILING DATE: 1998-06-09
APPLICATION NUMBER: 60/088722
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FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088861
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APPLICATION NUMBER: 60/089105
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APPLICATION NUMBER: 60/089512
FILING DATE: 1998-06-16
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                                                                  APPLICATION NUMBER: 60/087208
FILING DATE: 1938-05-28
APPLICATION NUMBER: 60/087609
FILING DATE: 1998-06-02
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APPLICATION NUMBER: 60/088028
FILING DATE: 1998-06-04
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FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088033
FILING DATE: 1998-06-04
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LICATION NUMBER: 60/088212
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FILING DATE: 1998-06-05
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NG DATE: 1998-06-10
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FILING DATE: 1998-06-10
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FILING DATE: 1998-06-10
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APPLICATION NUMBER: 60/088825
FILING DATE: 1998-06-10
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APPLICATION NUMBER: 60/088025
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FILING DATE: 1998-06-04
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FILING DATE: 1998-06-02
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FILING DATE: 1998-06-05
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Best Local Similarity 100.0
Matches 6; Conservative
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Gaps

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Indels

DB 14; Length 1089;

30.0%; Score 6; DB 14; Length 1089; 100.0%; Pred. No. 4.3e+02; Query Match Best Local Similarity

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PRIOR FILING DATE: 1997-10-24
PRIOR PLING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063121
PRIOR APPLICATION NUMBER: 60/06340
PRIOR PLING DATE: 1997-10-21
PRIOR PLING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-28
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PRIOR PRILING DATE: 1997-10-28
PRIOR PRILING DATE: 1997-10-28
PRIOR PRILING DATE: 1997-10-28
PRIOR PLING DATE: 1997-10-29
PRIOR PLING DATE: 1997-10-29
PRIOR PLING DATE: 1997-10-31
PRIOR PELING DATE: 1997-10-31
PRIOR PLING DATE: 1997-10-31
PRIOR PLING DATE: 1997-10-31
PRIOR PLING DATE: 1997-10-31
PRIOR PLING DATE: 1997-11-24
PRIOR PLING DATE: 1997-12-12
PRIOR PLING DATE: 1997-12-12
PRIOR PLING DATE: 1999-13-11
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PRIOR PLING DATE: 1999-13-11
PRIOR PLING DATE: 1999-3-11
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APPLICATION NUMBER: 60/082704
FILING DATE: 1998-04-22
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APPLICATION NUMBER: 60/080107
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PRIOR FILING DATE: 1998-03-31
PRIOR APPLICATION NUMBER: 60/080327
PRIOR FILING DATE: 1998-04-01
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PRIOR APPLICATION NUMBER: 60/081838
PRIOR FILING DATE: 1998-04-15
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PRIOR APPLICATION NUMBER: 60/081195
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APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Zhang, Zemin
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SCENETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C50
CURRENT EPLING DATE: 2022-06-19
CURRENT FILING DATE: 2022-06-19
CURRENT FILING DATE: 2022-06-19
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NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
  0; Mismatches
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CURRENT FILING DATE: 2002-06-18
PRIOR APPLICATION NUMBER: 10/02286
PRIOR FILING DATE: 2002-01-15
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059266
PRIOR PILING DATE: 1997-09-18
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Pred. No.
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100.0%; Pre
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Wood, William I.
Zhang, Zemin
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Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
  6; Conservative
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Smith, Victoria
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; ORGANISM: Homo Sapien
US-10-175-737-266
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US-10-175-737-266
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; Publication No. US200
; GENERAL INFORMATION:
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PRIOR FILING DATE: 1998-06-04
PRIOR PAPLICATION NUMBER: 60/088157
PRIOR APPLICATION NUMBER: 60/088167
PRIOR PILING DATE: 1998-06-05
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PRIOR PILING DATE: 1998-06-05
PRIOR PELING DATE: 1998-06-05
PRIOR PELING DATE: 1998-06-05
PRIOR PRING DATE: 1998-06-05
PRIOR PRING DATE: 1998-06-05
PRIOR PELING DATE: 1998-06-07
PRIOR PELING DATE: 1998-06-07
PRIOR APPLICATION NUMBER: 60/08855
PRIOR FILING DATE: 1998-06-09
PRIOR APPLICATION NUMBER: 60/088655
PRIOR PILING DATE: 1998-06-09
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APPLICATION NUMBER: 60/087208
FILING DATE: 1998-05-28
APPLICATION NUMBER: 60/087609
FILING DATE: 1998-06-02
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FILING DATE: 1998-04-29
APPLICATION NUMBER: 60/083559
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APPLICATION NUMBER: 60/084643
FILING DATE: 1998-05-07
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FILING DATE: 1998-05-15
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FILING DATE: 1998-05-15
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APPLICATION NUMBER: 60/086486
APPLING DATE: 1998-05-22
APPLICATION NUMBER: 60/097098
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APPLICATION NUMBER: 60/088025
FILING DATE: 1998-06-04
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FILLING DATE: 1998-06-04
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FILING DATE: 1998-06-04
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                                                                                                FILING DATE: 1998-04-29
APPLICATION NUMBER: 60/083496
FILING DATE: 1998-04-29
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FILING DATE: 1998-05-05
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APPLICATION NUMBER: 60/085579
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FILING DATE: 1998-05-15
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FILING DATE: 1998-06-02
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FILING DATE: 1998-06-04
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FILING DATE: 1998-05-06
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APPLICATION NUMBER: 60/082797
                                                          1998-04-28
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CURRENT FILING DATE: 2002-06-20
Prior application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
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PRIOR FILING DATE: 1998-06-10
PRIOR APPLICATION NUMBER: 60/08811
PRIOR FILING DATE: 1998-06-10
PRIOR APPLICATION NUMBER: 60/08824
PRIOR PELING DATE: 1998-06-10
PRIOR PERIOR PROPE: 1998-06-10
PRIOR PELING DATE: 1998-06-10
PRIOR FILING DATE: 1998-06-10
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PRIOR PELING DATE: 1998-06-11
PRIOR APPLICATION NUMBER: 60/08863
PRIOR PELING DATE: 1998-06-11
PRIOR PELING DATE: 1998-06-11
PRIOR APPLICATION NUMBER: 60/08909
PRIOR FILING DATE: 1998-06-12
PRIOR PELING DATE: 1998-06-16
PRIOR PELING DATE: 1998-06-17
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Gurney, Austin I.
Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Best Local Similarity 100.8
Matches 6, Conservative
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Goddard, Audrey
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ORGANISM: Homo Sapien
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GENERAL INFORMATION
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Gurney, Austin L.
Pan, James
Smith, Victoria
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Gurney, Austin L.
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Goddard, Audrey
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Best Local Similarity
                  12 WLRRYG 17
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US-10-173-706-266
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US-10-176-915-266
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SCRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C76
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430RIC83
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SEQ ID NO 266
LENGTH: 1089
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CURRENT FILING DATE: 2002-06-20
Prior application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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                                                                                   Sequence 266, Application US/10176749
Publication No. US20030017542A1
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Gurney, Austin L.
Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
Zhang, Zemin
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Watanabe, Colin K.
Wood, William I.
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Best Local Similarity 100.
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Goddard, Audrey
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; ORGANISM: Homo Sapien
US-10-176-749-266
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; ORGANISM: Homo Sapien
US-10-176-914-266
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687 WLRRYG 692
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                                                                                                                    GENERAL INFORMATION:
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LENGTH: 1089
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity luv...
6; Conservative
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Desnoyers, Luc
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ORGANISM: Homo Sapien
   ; ORGANISM: Homo Sapien
US-10-175-752-266
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US-10-176-482-266
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LENGTH: 1089
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APPLICANT: Goddord, Audrey
APPLICANT: Goddowski, Paul J.
APPLICANT: Goddowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Martin, Victoria
APPLICANT: Martanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Mood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
CURRENT APPLICATION NUMBER: US/10/175,738
CURRENT FILING DATE: 2002-06-19
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C60
CURRENY APPLICATION NUMBER: US/10/175,752
CURRENY FILING DATE: 2002-06-19
Frior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
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      30.0%; Score 6; DB 14; Length 1089; 100.0%; Pred. No. 4.3e+02; ive 0; Mismatches 0; Indels
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NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
                                                                                                                                                                                                                                                                                                         US-10-175-738-266

Sequence 266, Application US/10175738

Publication No. US20030022294A1

GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Goddard, Audrey
Godowski, Paul J
Gurney, Austin L
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Best Local Similarity 100.
Query Match
Best Local Similarity 100.
Matches 6; Conservative
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; ORGANISM: Homo Sapien
US-10-175-738-266
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APPLICANT:
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APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Pan, James
APPLICANT: Smith, Victoria
APPLICANT: Matanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: APPLICANTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430RIT70
CURRENT APPLICATION NUMBER: US/10/176,482
CURRENT FILING DATE: 2002-06-20
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERENCE: P3430R1C86
CURRENT APPLICATION NUMBER: US/10/176,757
CURRENT APPLICATION NUMBER: US/10/176,757
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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30.0%; Score 6; DB 14; Le
Best Local Similarity 100.0%; Pred. No. 4.3e+02;
Matches 6; Conservative 0; Mismatches 0;
                Score 6; DB 14; Login Pred. No. 4.3e+02;
30.0%; Scc. No. ...
100.0%; Pred. No. ...
0; Mismatches
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Publication No. US20030022297A1
GENERAL INFORMATION:
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Gurney, Austin L.
Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Gaps

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APPLICANT: Watenabe, Colin K.
APPLICANT: Watenabe, Colin K.
APPLICANT: Mood, William I.
APPLICANT: Abong, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P34.00RIC147
CURRENT APPLICATION NUMBER: US/10/180,557
CURRENT APPLICATION TEMOVED - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
SEQ ID NO S.612
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               Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612 SEQ ID NO 266 LENGTH: 1089
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APPLICANT: Baker Kevin P.
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Publication No. US20030022301A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Desnoyers, Luc
   CURRENT FILING DATE: 2002-06-25
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Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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Godowski, Paul J.
Gurney, Austin L.
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Smith, Victoria
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Matches 6; Conservative
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                                                                                           ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-180-552-266
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ORGANISM: Homo Sapien
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Best Local Similarity
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                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ITILE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                  Gaps
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                                                                                                        30.0%; Score 6; DB 14; Length 1089; 100.0%; Pred. No. 4.3e+02; ive 0; Mismatches 0; Indels
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CURRENT APPLICATION NUMBER: US/10/176,913
CURRENT FILING DATE: 2002-06-20
Prior Application removed - See file Wrapper or Palm NUMBER OF ESQ ID NOS: 612
SEQ ID NO 266
                                                                                                                                                                                                                                                                   RESULT 57
US-10-176-913-266
; Sequence 266, Application US/10176913
; Publication No. US20030022298A1
; GRNERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
appLICANT: Chen, Jian
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Smith, Victoria
Watanabe, Colin K.
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Goddard, Audrey
Godowski, Paul J.
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Gurney, Austin L.
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                                                                                                        Query Match
Best Local Similarity 100.
Matches 6; Conservative
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Goddard, Audrey
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                                    TYPE: PRT
CORGANISM: Homo Sapien
US-10-176-757-266
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Best Local Similarity
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US-10-180-552-266
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; SEQ ID NO 266
; LENGTH: 1089
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Watanabe, Colin K.
Wood, William I.
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Gurney, Austin L.
Pan, James
Smith, Victoria
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Matches 6; Conservative
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Goddard, Audrey
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ORGANISM: Homo Sapien
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US-10-174-582-266
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US-10-174-588-266
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APPLICANT: Chen, Jian
APPLICANT: Chen, Jian
APPLICANT: Gooddard, Audrey
APPLICANT: Goodwaki, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Fan, James
APPLICANT: Matanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: APPLICANT: ACOLIN K.
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SEQ ID VO 266
LENGTH: 1089
TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430R1014 CURRENT APPLICATION UNMBER: US/10/173,700 Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: PA30RLC40 CURRENT APLICATION NUMBER: US/10/174,572 CURRENT FILING DATE: 2002-06-19
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Publication No. US20030027264A1
GENERAL INFORMATION:
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Smith, Victoria
Watanabe, Colin K.
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Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
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ORGANISM: Homo Sapien
                                                                                                                                                          TYPE: PRT; ORGANISM: Homo Sapien
US-10-173-700-266
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                   SEQ ID NO 266
LENGTH: 1089
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APPLICANT:
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERENCE: P3430R1C36
CURRENT APPLICATION NUMBER: US/10/174,582
CURRENT FILING DATE: 2002-06-18
APPLICANT: Chang, Zenin Transmembrane PolyPepTides AND NUCLEIC TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: SECRETED AND THE SAME FILLE REFERENCE: P3430R1C31 CURRENT APPLICATION NUMBER: US/10/174,579 CURRENT FILING DATE: 2002-06-18 Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612 SEQ ID NO 266
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100.0%; Pred. No.
tive 0; Mismatch
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC FITLE OF INVENTION: ACIDS ENCODING THE SAME FILL REPERBNCE: P343 ORLGIS ENCODING THE SAME CURRENT PAPELICATION NUMBER: US/10/175,740

CURRENT APPLICATION NUMBER: US/10/175,740

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100.0%; Pred. No. 4.3e+02;
tive 0; Mismatches 0;
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PRIOR FILING DATE: 2002-01-15
PRIOR FILING DATE: 2002-01-15
PRIOR PAPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR PILING DATE: 1997-09-18
PRIOR PELING DATE: 1997-09-18
PRIOR PELING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-24
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PRIOR FILING DATE: 1997-10-21
PRIOR FILING DATE: 1997-10-21
PRIOR FILING DATE: 1997-10-28
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PRIOR FILING DATE: 1997-10-28
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Publication No. US20030027269A1
GENERAL INFORMATION:
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APPLICATION NUMBER: 60/063564
FILING DATE: 1997-10-28
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APPLICATION NUMBER: 60/063544
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Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Zhang, Zemin
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Smith, Victoria
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                                                                     Pan,James
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Best Local Similarity
Matches 6; Conserv
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US-10-175-743-266
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                                          APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TTLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLBIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P34.3081C28
CURRENT APPLICATION NUMBER: US/10/174,588
CURRENT APPLICATION NUMBER: US/2-06-18
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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CURRENT APPLICATION NUMBER: US/10/175,739
CURRENT FILING DATE: 2002-06-19
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100.0%; Pred. No. 4.3
ive 0; Mismatches
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Pred. No.
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o. US20030027267A1
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Publication No. US20030027268A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Desnoyers, Luc
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Wood, William I.
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Sequence 266, Application:
Sequence 266, Application No. US20030027267;
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
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Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
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Smith, Victoria
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ORGANISM: Homo Sapien
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ORGANISM: Homo Sapien
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687 WLRRYG 692
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LENGTH: 1089
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US-10-175-740-266
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LENGTH: 1089
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PRIOR FILING DATE: 1998-05-06

PRIOR APPLICATION UNMERS: 60/04640

PRIOR APPLICATION NUMBER: 60/04641

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PRIOR APPLICATION NUMBER: 60/04643

PRIOR APPLICATION NUMBER: 60/04551

PRIOR APPLICATION NUMBER: 60/05579

PRIOR PAPLICATION NUMBER: 60/05579

PRIOR PAPLICATION NUMBER: 60/06579

PRIOR FILING DATE: 1998-05-15

PRIOR PAPLICATION NUMBER: 60/06579

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PRIOR PAPLICATION NUMBER: 60/06632

PRIOR APPLICATION NUMBER: 60/06632

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PRIOR PAPLICATION NUMBER: 60/06632

PRIOR PAPLICATION NUMBER: 60/08632

PRIO

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APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430R1C107
CURRENT APPLICATION NUMBER: US/10/176,492
CURRENT FILING DATE: 2002-06-21
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APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEWERANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE OF INVENTION: ACIDS ENCODING THE SAME
FILE PREFERENCE: P4430R1692
CURRENT APPLICATION NUMBER: US/10/176,747
CURRENT APPLICATION NUMBER: US/10/176,747
FILING DATE: 2002-06-20
Prior application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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Mn. 4.38+02;
0; Indels
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NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
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100.0%; Pred. No. 4.3
tive 0; Mismatches
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GENERAL INFORMATION:
APPLICANT: Baker, Kevin P. APPLICANT: Chen, Jian
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Publication No. US20030027274A1
GRNERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Chen, Jian
APPLICANT: Gendyers, Luc
APPLICANT: Goddard, Audrey
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Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 100.
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Best Local Similarity
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US-10-176-750-266
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CURRENT APPLICATION NUMBER: US/10/176,488
CURRENT FILING DATE: 2002-06-21
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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; Sequence 266, Application US/10176492
; Publication No. US20030027272A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Chen, Jian
; APPLICANT: Goddard, Audrey;
; APPLICANT: Goddard, Audrey
PRIOR APPLICATION NUMBER: 60/089090
PRIOR FILING DATE: 1998-06-12
PRIOR PELICATION NUMBER: 60/089105
PRIOR PILING DATE: 1998-06-12
PRIOR PRING DATE: 1998-06-16
PRIOR PRING DATE: 1998-06-16
PRIOR PRING DATE: 1998-06-16
PRIOR PILING DATE: 1998-06-17
PRIOR PRING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089538
PRIOR PILING DATE: 1998-06-17
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Publication No. US20030027271A1
GENERAL INFORMATION:
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Gurney, Austin L.
Pan, James
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Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SCREETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C93
CURRENY APPLICATION NUMBER: US/10/176,987
CURRENY FILING DATE: 2002-06-21
PRIOR APPLICATION NUMBER: OS 06-21
PRIOR APPLICATION NUMBER: SECOND SECOND SECOND NOS: 612
FILE REFERENCE: PAGE OF SECOND NOS: 612
SECOND NO 266
LENGTH: 1089
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERENCE: P3430R1C100
CURRENT APPLICATION NUMBER: US/10/176,992
CURRENT APPLICATION NUMBER: US/10/176,992
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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Publication No. US20030027279A1
GENERAL INFORMATION:
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US-10-176-993-266
: Sequence 266, Application US/10176993
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Watanabe, Colin K.
Wood, William I.
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                                                        Godowski, Paul J.
Gurney, Austin L.
Pan, James
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Goddard, Audrey
Godowski, Paul J.
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                       Desnoyers, Luc
Goddard, Audrey
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ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo Sapien
    Chen, Jian
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                                                                                                                                  TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REPERBLICE: P3430R1C103
CURRENT APPLICATION NUMBER: US/10/176,750
CURRENT PILING DATE: 2002-06-21
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/176,985
CURRENT FILING DATE: 2002-06-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 266, Application US/10176985; Publication No. US20030027277A1; GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Goddard, Audrey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 266, Application US/10176987; Publication No. US20030027278A1; GENERAL INFORMATION: APPLICANT: Baker, Kevin P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
                                                      Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
Pan, James
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Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: P3430R1C99
                                                                                                                        Zhang, Zemin
                                                                                                                                                                                                                                                                                                                        TYPE: PRT
CORGANISM: Homo Sapien
US-10-176-750-266
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LENGTH: 1089
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## SPELICANT: Baker, Kevin P.
## APPLICANT: Baker, Kevin P.
## APPLICANT: Chen, Jian
## APPLICANT: Chen, Jian
## APPLICANT: Goddard, Audrey, Jusc
## APPLICANT: Goddard, Audrey, Jusc
## APPLICANT: Goddard, Audrey, Jusc
## APPLICANT: Goddard, Audrey, Austin L.
## APPLICANT: Godrey, Austin L.
## APPLICANT: Ban, James
## APPLICANT: Ban, James
## APPLICANT: Smith, Victoria
## APPLICANT: Smith, Indepression
## APPLICANT: Smith, Indepression
## APPLICANT: The APPLICANT: Smith, Indepression
## APPLICANT: Smith, Indepression
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Search completed: October 26, 2004, 07:47:54 Job time : 66 secs

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Sequence 3148, A
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Sequence 22818, A
Sequence 17567, A
Sequence 7940, Ap
Sequence 7940, Ap
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Sequence 20409, A
Sequence 20409, A
Sequence 17629, A
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2: /cgn2_6/ptcdata/1/iaa/5B_COMB.pep:*

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4: /cgn2_6/ptcdata/1/iaa/6B_COMB.pep:*

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   version 5.1.6
- 2004 Compugen Ltd.
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US-09-27-122
US-09-27-291A-31470
US-09-270-76-31648
US-09-270-76-46865
US-09-270-76-46865
US-09-252-991A-17667
US-09-252-991A-17667
US-09-252-991A-17667
US-09-252-991A-17667
US-09-252-991A-2019
US-09-252-991A-2019
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US-09-65-270A-3
US-09-65-270A-3
US-09-65-270A-3
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US-09-65-270A-3
US-09-98-024C-1
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                       protein search, using sw model
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ADDRESSEE: Fish & Richardson, P.C.
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APPLICANT: McCoy, John M.
APPLICANT: Jessen, Timm H.
APPLICANT: Xu, Chanxing Wilson
TITLE OF INVENTION: INVERACTION TRAP SYSTEMS FOR DETECTING PROTEIN
NUMBER OF SEQUENCES: 28
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                                                                                                                               Sequence 12, Application US/08504538A

Patent No. 600446

GENERAL INFORMATION:
APPLICANT: Brent, Roger
APPLICANT: McCoy, John M.
APPLICANT: Dessen, Timm H.
TITLE OF INVENTION: PROTEIN INTERACTION FRAP SYSTEMS FOR DETECTING TITLE OF INVENTION: PROTEIN INTERACTIONS

NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESSE: 21
STREET: 176 Federal Street
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/504,538A
FILING DATE: 07/20/95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 20; DB 3; L. ilarity 100.0%; Pred. No. 1.4e-14; Conservative 0; Mismatches 0;
                                              ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 00786/259001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 428-0200
TELEPAX: (617) 428-045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-630-052-12; Sequence 12, Application US/08630052; Patent No. 6399296; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08/278,082
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20
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Matches 20, Conserv
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ZIP: 02110-2214
                                                                                                                                                                                                                                                                                                                                                                                         Boston
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CORRESPONDENCE ADDRESS:

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GENERAL INFORMATION:
APPLICANT: Brent, Roger
APPLICANT: McCoy, John M.
APPLICANT: Jessen, Timm H.
APPLICANT: Jessen, Timm H.
APPLICANT: V., Chanxing Wilson
TITLE OF INVENTION: DETECTING PROTEIN INTERACTIONS
                                                                      COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 20; DB 3; I
100.0%; Pred. No. 1.4e-14;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
COMFUTER: IBM PC compatible
COMFUTER: BATENT PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
SOFTWARE: #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/09307
FILING DATE:
CLASSIFICATION:
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3R: 00786/311001
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225 Franklin Street
                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/504,538
FILING DATE: July 20, 1995
APPLICATION NUMBER: 08/278,082
FILING DATE: July 20, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Karen F. Lech
REGISTRATION NUMBER: 35,238
REFERENCE/DOCKET NUMBER: 00786/31
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                     Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
US-08-630-052-12
ALLEET: STREET: TW: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCT-US95-09307-12
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TELEX: 20
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Indels

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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: ABRCGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196,136

CURRENT APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR FILING DATE: 1998-07-18

PRIOR FILING DATE: 1998-07-27

NUMBER: OS EQ ID NOS: 33142

SEQ ID NO 32834

LENGTH: 138
                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:

APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE PEFERENCE: File Reference: 7326-094
CURRENT APPLICATION NURBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 46665
LENGTH: 135
                                                                                               Length 135;
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                                                                                            DB 4;
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30.0%; Score 6; DB 4;
Best Local Similarity 100.0%; Pred. No. 24;
Matches 6; Conservative 0; Mismatches
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                                                                                        30.0%; Score 6; DB 4
100.0%; Pred. No. 24;
tive 0; Mismatches
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; Patent No. 6551795
                                                                                                                                                                                                                                                                                                                       Sequence 46865, Application US/09270767 Patent No. 6703491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30.0%; SCC.
100.0%; Pre
        TYPE: PRT
ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Drosophila melanogaster US-09-270-767-46865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                   6; Conservative
                                                                                          Query Match
Best Local Similarity
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                                               US-09-270-767-31648
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APPLICANT: MARC J. Rubenfield et al.
APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERBNES: 107196,136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,190
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEGO ID NO 31470
LENGTH: 480
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TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILLE REFERENCE: FILE REFERENCE: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 31648
LENGTH: 135
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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100.0%; Pred. No. 0.5
ative 0; Mismatches
                                                      ATTORNAY AGENT INFORMATION:
NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/288001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 31470, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-270-767-31648
; Sequence 31648, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
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                                                                                                                                                                                   TELEFAX: (617) 542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100."
Matches 20, Conservative
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Best Local Similarity 100.
Matches 8; Conservative
PRIOR APPLICATION DATA:
                   APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                                                                                               amino acid
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                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANT TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANTITLE OF INVENTION: POR DIAGNOSTICS AND THERAPEUTICS
TITLE REPRESENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 20409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5410, Application US/09134000C

Sequence 5410, Application US/09134000C

Patent No. 6617156

GREERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

TITLE OF INVENTION: ENTEROCCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REPERBENCE: 03296-032

CURRENT APPLICATION NUMBER: US/09/134,000C

CURRENT PILING DATE: 1998-08-13

PRIOR FILING DATE: 1998-08-15

NUMBER OF SEQ ID NOS: 6812

SOFTWARE: Patentin version 3.1
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                                                                                                                                                          30.0%; Score 6; DB 4;
100.0%; Pred. No. 56;
tive 0; Mismatches
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100.0%; Pred. No. 59;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                      RESULT 11
US-00-248-796A-20409
; Sequence 20409, Application US/09248796A
; Patent No. 6747137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enterococcus faecalis
                                                                                      ORGANISM: Klebsiella pneumoniae
US-09-489-039A-7940
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 7940
LENGTH: 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Candida albicans
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Best Local Similarity 100...
...a 6; Conservative
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                                                                                                                                                                                                            6; Conservative
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                   388 WLRRYG 393
                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                          12 WLRRYG 17
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LENGTH: 439
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                                                                     TYPE: PRT
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                               APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERGGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196_136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
SEQ ID NO 26818
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 26818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

APPLICANT: Marc J. Rubenfield et al.

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: AERGGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196,136

CURRENT APPLICATION NUMBER: US 60/074,788

PRIOR PILING DATE: 1999-02-18

PRIOR PILING DATE: 1999-02-18

PRIOR PILING DATE: 1999-07-27

NUMBER: OS 60/094,190

RIOR PILING DATE: 1999-07-27

NUMBER: OS 33142

SEQ 1D NO 17567

LENGTH: 277
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Sequence 7940, Application US/08489039A

Patent No. 6610836

GENERAL INFORMATION:

APPLICANT:

APPLICANT:

TITLE OF INVENTION:

PITLE OF INVENTION:

TITLE OF INVENTION:

TITLE OF INVENTION:

PREDMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
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100.0%; Pred. No. 29;
tive 0; Mismatches
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100.0%; Pred. No. 42,
:ive 0; Mismatches
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CURRENT APPLICATION NUMBER: US/09/489,039A

CURRENT FILING DATE: 2000-01-27

PRIOR PELING DATE: 1999-01-29

PRIOR FILING DATE: 1999-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17567, Application US/09252991A Patent No. 6551795
  Sequence 26818, Application US/09252991A
Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-26818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-17567
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Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity
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Gaps
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Sequence 16, Application US/09370700
Sequence 16, Application US/09370700
Sequence 17, Richard H
APPLICANT: Baltz, Richard H
APPLICANT: Crawicord, Kathryn P
APPLICANT: Treadway, Patti J
APPLICANT: Turner, Jan R
APPLICANT: Turner, Jan R
APPLICANT: Waldron, Clive
ITILE OF INVENTION: Biosynthetic Genes For Spinosyn Insecticide
ITILE OF INVENTION: Biosynthetic Genes For Spinosyn Insecticide
SIERERRY APPLICATION NUMBER: US/09/370,700
CURRENT FILING DATE: 1999-08-09
CURRENT FILING DATE: 1999-08-09
SADFIRER FILING DATE: 1999-03-09
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn Ver. 2.0
                                           APPLICANT: Treadway, Patti J.

APPLICANT: Turner, Jan R.

APPLICANT: Waldron, Clive
TITLE OF INVENTION: Biosynthetic Genes For Spinosyn Insecticide
TITLE OF INVENTION: Production
NUMBER OF SEQUENCES: 39
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                  ZIP: 4626B

COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/036,987A
FILING DATE: 09-MAR-1998
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: STUART, DONAIG R
REFERENCE/DOCKET NUMBER: 28,479
REFERENCE/DOCKET NUMBER: 50,608
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
                                                                                                                                                STREE: DOW AGYOSCIENCES LLC Patent Department STREET: 9330 Zionsville Road CITY: Indianapolis
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                        Merlo, Donald J.
Treadway, Patti J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (317)337-4847
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100.00
The 6; Conservative
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amino acid
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US-09-370-700-16
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                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
FILE REPERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 17629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MARC J. Rubenfield et al.
APPLICANT:
APPLICANT:
MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS;
TITLE OF INVENTION: AREUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS.
TITLE OF INVENTION: APEUCINOSA FOR DIAGNOSTICS AND THERAPEUTICS.
FILE REFERENCE: 107196,136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR PLING DATE: 1999-02-18
PRIOR FILING DATE: 1999-07-27
NUMBER OF SEQ ID NOS: 33142
SEGO ID NO 22078
LENGTH: 478
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100.0%; Pred. No. 61;
tive 0; Mismatches
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0; Mismatches
                                                                                      US-09-252-991A-17629
; Sequence 17629, Application US/09252991A
; Patent No. 6551795
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Sequence 16, Application US/09036987A
Patent No. 6143526
GENERAL INFORMATION:
APPLICANT: Baltz, Richard H.
APPLICANT: Broughton, Mary C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-17629
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ORGANISM: Pseudomonas aeruginosa
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Best Local Similarity 100.
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Matches 6; Conservative
193 SLWALG 198
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Query Match
Best Local Similarity 10v...
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Matches 6; Conserv
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                                                                     Patent No. 6329151
GENERAL INFORMATION:
                               US-09-655-270A-3
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APPLICANT:

APPLICANT:

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA

CURRENT PER EXPERIENCE: 2709-2004001

CURRENT FILING NOTE: 2000-01-27

PRIOR APPLICATION NUMBER: US 60/117,747

PRIOR PLILING DATE: 1999-01-29

NUMBER OF SEQ ID NOS: 14342

SEQ ID NO 8255
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                                                                                                                                                                                                                                      Parent No. 6221406
GRNERAL INFORMATION:
APPLICANT: Baltz, Richard H
APPLICANT: Baltz, Richard H
APPLICANT: Crawford, Kathryn P
APPLICANT: Madouri, Krishnamurthy
APPLICANT: Turner, Jan R
APPLICANT: Turner, Jan R
APPLICANT: Turner, Jan R
APPLICANT: Turner, Jan R
APPLICANT: Waldron, Clive
TITLE OF INVENTION: Biosythetic Genes For Spinosyn Insecticide
FILE REFERENCE: 50489 DIV1
CURRENT APPLICATION NUMBER: US/09/603,207B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 486;
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             DB 3; Length 486; . 66;
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30.0%; Scor.
100.0%; Pred. No. ...
0; Mismatches
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100.0%; Pred. No. 66;
iive 0; Mismatches
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100.0%; Pred. No. 67;
cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2000-06-23
BARLIER APPLICATION NUMBER: 09/370,700
BERLIER FILING DATE: 1998-03-09
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PATCHIEN Ver. 2.0
SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8255, Application US/09489039A Patent No. 6610836
                                                                                                                                                                                                  US-09-603-207-16
; Sequence 16, Application US/09603207B
: Patent No. 6521406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ) ORGANISM: Saccharopolyspora spinosa US-09-603-207-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-8255
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Matches 6; Conservative
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Best Local Similarity 100.
Matches 6; Conservative
                                                     6; Conservative
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               Query Match
Best Local Similarity
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APPLICANT: Rouviere, Pierre E.

TITLE OF INVENTION: High Density Sampling of Differentially Expressed Prokaryotic mRI FILE REPERENCE: BC1001 US NA CURRENT APPLICATION NUMBER: US/09/655,270A CURRENT APPLICATION NUMBER: 06/120,702 PRIOR APPLICATION NUMBER: 06/120,702 PRIOR APPLICATION NUMBER: 60/120,702 PRIOR APPLICATION NUMBER: 60/125,542 PRIOR FILING DATE: 1999-September-03 NUMBER OF SEQ ID NOS: 37 SOFFWARE: Microsoft Office 97 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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CURRENT APPLICATION NUMBER: US/09/651,941
CURRENT APPLICATION NUMBER: US/09/651,941
CURRENT-FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/152,545
PRIOR APPLICATION NUMBER: 60/152,545
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Microsoft Office 97
Lawrent Company No 10 No 1
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Sequence 3, Application US/09955597

Patent No. 6441856

GENERAL INFORMATION:

APPLICANT: WALTER, DIRRE E

APPLICANT: RAINER, RUSS

TITLE OF INVENTION: Genes Encoding Picric Acid Degradation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 532;
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100.0%; Pred. No. 71;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 6; DB 3;
Pred. No. 71;
0; Mismatches
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Patent No. 6355470
GENERAL INFORMATION:
APPLICANT: ROUVIER, PIERRE E
APPLICANT: RAINER, RUSS
APPLICANT: RAINER, RUSS
TITLE OF INVENTION: Genes Encoding P
FILE REFERENCE: BC1022 US NA
Sequence 3, Application US/09655270A Patent No. 6329151
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100.0%; Pr
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Patent No. 5719064
GENERAL INFORMATION:
APPLICANT: Socield, R. Hal
APPLICANT: Socield, R. Hal
APPLICANT: Harley, John B.
TITLE OF INVENTION: Spendyloarthropathies
UNMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
                                                                                                                                                                                              0; Indels
                                                                                                                                                      DB 4; Length 12; .37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Atlanta
CITY: Atlanta
CITY: Atlanta
COUNTRY: USA
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHIT RE-BEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/944,143C
FILING DATE:
CLASSIFICATION: 436
                                                                                                                                                 25.0%; Score 5; DB 4
100.0%; Pred. No. 37;
ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 25.0%; Score 5; DB 1
Best Local Similarity 100.0%; Pred. No. 54;
Matches 5; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OMRF138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: Pablst, Pattrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMRE
TELECOMMUNICATION:
TELEPHONE: (404)873-8794
                                                                 FEATURE:
; OTHER INFORMATION: Oligopeptide
US-08-988-024C-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (404) 873-8795
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 19 amino acids amino acids
                                                                                                                                                                                              5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                 Query Match
Best Local Similarity
Matches 5; Conserv
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N
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            LENGTH: 12
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APPLICANT: Nacc J. Rubenfield et al.
APPLICANT: Nacc J. Rubenfield AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
FILES REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 24635
LENGTH: 800
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APPLICANT: Monfeotre, Joseph A.
APPLICANT: Becker, Christopher H.
APPLICANT: Becker, Daniel J.
APPLICANT: Sollart, Daniel J.
TITLE OF INVENTION: Releasable No. 6635452volatile Mass-Label Molecules
FILE REFERENCE: 24736-2057
                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                    30.0%; Score 6; DB 4;
100.0%; Pred. No. 71;
tive 0; Mismatches
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100.0%; Pred. No. 98;
tive 0; Mismatches
FILE REFERENCE: BC1022 US NA
CURRENT APPLICATION NUMBER: US/09/955,597
CURRENT FILING DATE: 2001-09-17
FRIOR APPLICATION NUMBER: 60/152,545
FRIOR FILING DATE: 1999-10-03
NUMBER OF SEQ ID NOS: 28
SOTWARE: Microsoft Office 97
SEQ ID NO 3
LENTH: 532
                                                                                                                                                                                   ; TYPE: PRT
; ORGANISM: Rhodococcus erythropolis HL PM-1
US-09-955-597-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24635, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 7, Application US/08988024C Patent No. 6635452
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ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 6; Conservative
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Best Local Similarity
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US-09-252-991A-24635
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US-08-988-024C-7
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Gaps
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IS-07-944-143C-20
Sequence 20. Application US/07944143C
Sequence 20. Application US/07944143C
Sequence 20. Application US/07944143C
Sequence 20. Application
TITORNI Socield, R. Hal
APPLICANT: Socield, R. Hal
TITLE OF INVENTION: Peptide Diagnostics and Therapeutics for TITLE OF INVENTION: Spondyloarthropathies
ITLE OF INVENTION: Spondyloarthropathies
NUMBER OF SEQUENCES:
NUMBER OF SEQUENCES:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 2800 One Atlantic Center
STREET: 2800 Usest Peachtree Street
STREET: Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: amino acid sequence of the icosapeptide MH20; OTHER INFORMATION: the extracellular virus-binding domain of the OTHER INFORMATION: second artificial receptor
                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Curiel, David T.
APPLICANT: Douglas, Joanne T.
APPLICANT: Krasnykh, Victor N.
APPLICANT: Krasnykh, Victor N.
APPLICANT: Frasnykh, Igor
ITTLE OF INVENTION: Fiber Receptor-Independent System for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                   0; Indels
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                                                                                                                                                                       Length 20;
                                                                                                                                                                         DB 4;
. 56;
                                                                                                                                                            25.0%; Scor.
100.0%; Pred. No. 50,
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: D6159
CURRENT APPLICATION NUMBER: US/09/498,134A
CURRENT FILING DATE: 2000-02-03
PRIOR APPLICATION NUMBER: US 60/118,880
PRIOR FILING DATE: 1999-02-05
NUMBER OF SEQ ID NOS: 11
SEQ ID NO 3
LENGTH: 20
                                                                                    ; FEATURE:
; OTHER INFORMATION: Synthetic peptide MH20
US-09-155-613A-6
                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 3, Application US/09498134A; Patent No. 6649396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: artificial sequence
                                         TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5; Conservative
                                                                                                                                                                                                                     Conservative
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Best Local Similarity
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Best Local Similarity
Matches 5; Conserv
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SEQ ID NO 6
LENGTH: 20
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                                                                                                                for Spondyloarthropat
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APPLICANT: Boulanger, Pierre
APPLICANT: Hong, Saw See
APPLICANT: Arayan, Lucie
ITILE OF INTENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
ITILE OF INTENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
ITILE OF INTENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
ITILE OF INTENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
ITILE OF INTENTION: US/09/155,613A
CURRENT FILING DATE: 1998-01-30
PRIOR FILING DATE: 1998-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-09-09
INUMBER OF SEQ ID NOS: 98
SOFTWARE: PastSEQ for Windows Version 4.0
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                                       Sequence 21, Application PC/TUS9308214
GENERAL INFORMATION:
APPLICANT: Oklahoma Medical Research Foundation
TITLE OF INVENTION: Peptide Diagnostics and Therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
                                                                                                                                                                                                                     CITY: Atlanta
STATE: Georgia
COUNTRY: United States
ZIP: 330309-4530
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08214
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25.0%; Scorr
100.0%; Pred. No. 5.,
                                                                                                                                                                              ADDRESSEE: Kilpatrick & Cody
STREET: 1100 Peachtree Street, Suite 2800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OMRF138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6, Application US/09155613A Patent No. 6420120 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
ATTORNEX/AGENT INFORMATION:
TAMBE: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REPRENCE/DOCKET NUMBER: OMRE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 815-6508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (404) 815-6555
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 25.(
Best Local Similarity 100.
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM:
    RESULT 25
PCT-US93-08214-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-155-613A-6
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GENERAL INFORMATION:

APPLICANT: Boulanger, Pierre

APPLICANT: Boulanger, Pierre

APPLICANT: Boulanger, Pierre

APPLICANT: Hong, Saw See

APPLICANT: Hong, Saw See

APPLICANT: Hardy Lucie

TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses

TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses

TITLE OF INVENTION NUMBER: US/09/155,613A

CURRENT FILING DATE: 1998-09-30

PRIOR FILING DATE: 1998-01-30

PRIOR PILING DATE: 1997-01-30

PRIOR APPLICATION NUMBER: FR 97/11166

PRIOR FILING DATE: 1997-09-09

NUMBER OF SEQ ID NOS: 98

SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Boulanger, Pierre
APPLICANT: Boulanger, Pierre
APPLICANT: Hong, Saw See
APPLICANT: Hong, Saw See
APPLICANT: Hong, Saw See
TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
FILE REPERBENCE: 032751-036
CURRENT APPLICATION NUMBER: US/09/155,613A
CURRENT FILING DATE: 1998-09-30
PRIOR FILING DATE: 1998-09-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR APPLICATION NUMBER: FR 97/11166
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-09-09
                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                           Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 24;
                                                                                                                                                                                      25.0%; Score 5; DB 5;
100.0%; Pred. No. 58;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25.0%; Score 5; DB 4 ilarity 100.0%; Pred. No. 65; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 97, Application US/09155613A Patent No. 6420120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/09155613A Patent No. 6420120 GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Phagotope US-09-155-613A-97
                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                         5; Conservative
                                                                           internal
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Best Local Similarity
The 5; Conserve
                                                                                                                                                                                                                 Best_Local Similarity
Matches 5; Conserv
  MOLECULE TYPE: pe
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: ir
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                     12 WLRRY 16
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                                                                                                                                             PCT-US93-08214-20
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US-09-155-613A-97
                                                                                                                    ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 31
US-09-155-613A-1
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                                                                                                                                                                                           Query Match
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GENERAL INFORMATION:
APPLICANT: Oklahoma Medical Research Foundation
TITLE OF INVENTION: Peptide Diagnostics and Therapeutics
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kilpatrick & Cody
STREET: 1100 Peachtree Street, Suite 2800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25.0%; Score 5; DB 1; Length 21; 100.0%; Pred. No. 58; 0; Indels tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Atlanta
STATE: Georgia
COUTRY: United States
ZIP: 310309-4530
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                 OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/944,143C
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PELLING DATE:
CLASSIPICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMF138
TELECOMMUNICATION INFORMATION:
TELECHONE: (404) 815-6508
TELECHONE: (404) 815-6508
INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acids
                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: PADEL, PATER 1, 284
REFERENCE/DOCKET NUMBER: OMRETEECOMUNICATION INFORMATION:
TELEPHONE: (404)873-8794
TELEPHONE: (404)873-8794
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                          21 amino acids
                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        internal
                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3S: single
linear
                                                                                                               FILING DATE:
CLASSIFICATION: 436
                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12 WLRRY 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO FRAGMENT TYPE: i ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 WLRRY 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-07-944-143C-20
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Gaps

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GENERAL INFORMATION:

APPLICANT: Boulanger, Pierre

APPLICANT: Boulanger, Lucie

APPLICANT: Boulanger, Lucie

APPLICANT: Boulanger, Lucie

TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses;
FILE REFERENCE: 037751-036

CURRENT APPLICATION NUMBER: US/09/155,613A

CURRENT FILING DATE: 1998-00-30

PRIOR APPLICATION NUMBER: PCT/FR98/00184

PRIOR APPLICATION NUMBER: FR 97/01005

PRIOR PILING DATE: 1997-01-30

PRIOR PILING DATE: 1997-01-30

PRIOR FILING DATE: 1997-09-09

NUMBER OF SEQ ID NOS: 98

SOFTWARE: FastSEQ for Windows Version 4.0
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.88;
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GENERAL TRUBEN et al.
TITLE OF INVENTION: 94 Human Secreted Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25.0%; Scc.
100.0%; Pred. No. c.
0; Mismatches
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EARLIER APPLICATION NUMBER: PCT/US99/13418

EARLIER APPLICATION NUMBER: 60/089,507

EARLIER FILING DATE: 1998-06-16

EARLIER FILING DATE: 1998-06-16
                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                        25.0%; Score 5; Discore 5; Discor
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EARLIER APPLICATION NUMBER: 60/090,112
EARLIER FILING DATE: 1998-06-22
EARLIER APPLICATION NUMBER: 60/090,113
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US-09-155-613A-23
Gequence 23, Application US/09155613A
Patent No. 6420120
                                                                                                                                                                                                            Query Match 25.0
Best Local Similarity 100.
Matches 5; Conservative
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; SEQ ID NO 22
; LENGTH: 35
; TYPE: PRT
; ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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US-09-461-325-242
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                                                                                                                ; OKGANALE... US-09-155-613A-22
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LENGTH: 35
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APPLICANT: Boulanger, Pierre
APPLICANT: Hong, Saw See
APPLICANT: Hong, Saw See
ITILE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses
FILE REPERENCE: 032751-036
CURRENT APPLICATION NUMBER: US/09/155,613A
CURRENT FILING DATE: 1998-09-30
FRIOR APPLICATION NUMBER: PCT/FR98/00184
FRIOR APPLICATION NUMBER: FR 97/01005
FRIOR APPLICATION NUMBER: FR 97/11166
FRIOR FILING DATE: 1997-09-09
NUMBER OF SEQ ID NOS: 98
NUMBER OF SEQ ID NOS: 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Boulanger, Pierre
APPLICANT: Hong, Saw See
APPLICANT: Hong, Saw See
APPLICANT: Arayan, Lucie
TITLE OF INVENTION: Use of a Polypeptide as Cell Receptor for Adenoviruses;
TITLE REFERENCE: 032751-036
CURRENT APPLICATION NUMBER: US/09/155,613A
CURRENT PILING DATE: 1998-09-30
PRIOR PRILING DATE: 1998-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
NUMBER OF SEQ ID NOS: 98
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                               Length 25;
                                                                                                                                                                                                                                            25.0%; Score 5; DB 4;
100.0%; Pred. No. 67;
iive 0; Mismatches
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100.0%; Pred. No. 67;
iive 0; Mismatches
       SOFTWARE: FastSEQ for Windows Version 4.0
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US-09-155-613A-98
; Sequence 98, Application US/09155613A
natent No. 6420120
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'Sequence 22', Application US/09155613A

'Patent No. 6420120

; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; FEATURE:
; OTHER INFORMATION: Phagotope
US-09-155-613A-98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 25.0
Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                      5; Conservative
                                                                                                             TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 WLRRY 16
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                                                                                                                                                       ; UKGAN15::...US-09-155-613A-1
                                          SEQ ID NO 1
LENGTH: 25
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GENERAL INFORMATION:

APPLICANT: Ruben et al.
TITLE OF INVENTION:
CURRENT REPERENCE: 94 Human Secreted Proteins
FILE OF INVENTION: 94 Human Secreted Proteins
FILE OF INVENTION: 94 Human Secreted Proteins
FILE OF INVENTION NUMBER: US/10/012,542
CURRENT FILING DATE: 2001-12-12
PRIOR APPLICATION NUMBER: EARLIER PELING DATE: 1999-12-14
PRIOR FILING DATE: EARLIER FILING DATE: 1999-16-16
PRIOR PELING DATE: EARLIER FILING DATE: 1998-06-16
PRIOR PELING DATE: EARLIER FILING DATE: 1998-06-22
PRIOR PELING DATE: EARLIER FILING DATE: 1998-06-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-22
NUMBER OF SEQ ID NOS: 532
SOFTWARE: PATENTIN VOWER: 2.0
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/089,508
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/089,509
PRIOR FILING DATE: BARLIER FILING DATE: 1998-06-16
PRIOR FILING DATE: BARLIER FILING DATE: 1998-06-16
PRIOR FILING DATE: BARLIER FILING DATE: 1998-06-16
PRIOR PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/090,112
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-22
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-22
NUMBER: OF SEQ ID NOS: 532
SOFTWARE: PACHLER FILING DATE: 1998-06-22
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100.0%; Pred. No. 92;
ative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 429, Application US/10012542 Patent No. 6627741
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; Patent No. 6774216
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 5; Conservative
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US-10-012-542-242
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Best Local Similarity
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US-10-115-123-242
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LENGTH: 37
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CURRENT FILING DATE: 2001-12-12
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/461,325
PRIOR FILING DATE: EARLIER FILING DATE: 1999-12-14
PRIOR PLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/089,507
PRIOR FILING DATE: EARLIER FILING DATE: 1998-06-16
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Patent No. 6627741
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 94 Human Secreted Proteins
FILE REFERENCE: PZ029P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 94 Human Secreted Proteins
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CURRENT FILING DATE: 1999-12-14
EARLIER APPLICATION NUMBER: UF/US99/13418
EARLIER FILING DATE: 1999-06-15
EARLIER FILING DATE: 1999-06-15
EARLIER FILING DATE: 1998-06-16
EARLIER APPLICATION NUMBER: 60/089,508
EARLIER APPLICATION NUMBER: 60/089,508
EARLIER FILING DATE: 1998-06-16
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EARLIER FILING DATE: 1998-06-12
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Pred. No.
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100.0%;
  EARLIER FILING DATE: 1998-06-22
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SEQ ID NO 429
                            NUMBER OF SEQ ID NOS: 532
SOFTWARE: Patentin Ver. 2.0
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Best Local Similarity 100.
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; ORGANISM: Homo sapiens
US-09-461-325-429
                                                                                                                           TYPE: PRT
CORGANISM: Homo sapiens
US-09-461-325-242
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Best Local Similarity
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                                                                         SEQ ID NO 242
LENGTH: 37
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Patent No. 5717061
GENERAL INFORMATION:
APPLICANT: Rao, Gururaj A.
APPLICANT: Zhong, Lingxiu
TITLE OF INVENTION: SYMTHETIC ANTIMICROBIAL PEPTIDES
NUMBER OF SEQUENCES:
ADDRESSEE: ADDRESSE:
ADDRESSEE: PIONEER HI-BRED INTERNATIONAL, INC.
                                                                                                                                                                                                              APPLICANT: Rao, A. Gururaj; Zhong, Lingxiu
TITLE OF INVENTION: SYNTHETIC ANTIMICROBIAL PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MIDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage COMPUTER: IBM Compatible OPERATING SYSTEM: MS-DOS/Microsoft Windows SOFTWARE: Microsoft Windows No. 5607914epad CURRENT APPLICATION NUMBER: US/08/179,632

FILING DATE: 07-JAN-1994
                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSER: Pioneer Hi-Bred International, Inc ATREET: 700 Capital Square, 400 Locust Street CITY: Des Moines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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100.0%; Pred. No.
:ive 0; Mismatch
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 07-JAN-1994
CLASSIFICATION: 530
PRICK APPLICATION DATE: 08/079,512
APPLICATION NUMBER: 08/079,512
FILING DATE: 06/18/93
ATTORNEY/AGENT INFORMATION:
NAME: ROCH, Michael J.
REGISTRATION NUMBER: 29,342
REFREENCE/DOCKET NUMBER: 0233 US
TELECOMMUNICATION INFORMATION:
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                                                                                                                                         Sequence 7, Application US/08179632; Patent No. 5607914; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                        STATE: Iowa
COUNTRY: United States
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                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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STRANDEDNESS: siz
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Matches 5; Conserv
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                              23 SLWAL 27
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US-08-440-174A-7
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APPLICANT: Ruben et al.
TITLE OF INVENTION: 94 Human Secreted Proteins
FILE REPRENCE: PSZ03G30ADP1D2
CURRENT PREPENCE: PSZ03G30ADP1D2
CURRENT APPLICATION NUMBER: US/10/115,123
CURRENT PILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: PCT/US99/13418
PRIOR FILING DATE: 1999-06-15
PRIOR PELING DATE: 1998-06-16
PRIOR PRILING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/089,509
PRIOR APPLICATION NUMBER: 60/089,509
PRIOR PLING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/089,510
PRIOR APPLICATION NUMBER: 60/089,510
PRIOR APPLICATION NUMBER: 60/089,510
PRIOR APPLICATION NUMBER: 60/089,112
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APPLICANT: Ruben et al.
TITLE OF INVENTION: 94 Human Secreted Proteins
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100.0%; Pred. No. 92;
live 0; Mismatches
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CURRENT APPLICATION NUMBER: US/10/115,123
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: PCT/US99/13418
PRIOR FILING DATE: 1999-06-15
PRIOR FILING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/090,112
PRIOR FILING DATE: 1998-06-16
PRIOR FILING DATE: 1998-06-16
PRIOR FILING DATE: 1998-06-17
PRIOR FILING DATE: 1998-06-22
PRIOR FILING DATE: 1998-06-22
NUMBER: OF SEQ. DE NOS: 532
                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1998-06-22
PRIOR APPLICATION NUMBER: 60/090,113
PRIOR FILING DATE: 1998-06-22
NUMBER OF SEQ ID NOS: 532
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 242
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Sequence 429, Application US/10115123
Patent No. 6774216
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SEQ ID NO 429
LENGTH: 37
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Best Local Similarity 100
Matches 5; Conservative
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CRGANISM: Homo sapiens
US-10-115-123-429
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 5; Conserv
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APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
NUMBER OF SEQUENCES: 7310
                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 25.0%; Score 5; DB 4; Length 60; Best Local Similarity 100.0%; Pred. No. 1.38+02; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                             Length 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085,598
FILING DATE: 14 MAY 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                DB 5;
                                                                                                                                                                             25.0%; Score 5; DB 5
100.0%; Pred. No. 94;
cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (B) LOCATION 1...60
SEQUENCE DESCRIPTION: SEQ ID NO: 4928:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
                                                                                      MOLECULE TYPE: protein
DESCRIPTION: hordothionin derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4928, Application US/09107532A Patent No. 6583275 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: PC
OPERATING SYSTEM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGIH: 60 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (781)893-8:
INFORMATION FOR SEQ ID NO: 493
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
                                                                                                                                                                                                                        5; Conservative
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                                                   single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                               amino acid
                                                                     linear
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                          TYPE: amino s
STRANDEDNESS:
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US-09-107-532A-4928
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                                                                     TOPOLOGY:
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        LENGIH:
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TITLE OF INVENTION: SYNTHETIC ANTIMICROBIAL PEPTIDES
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pioneer Hi-Bred International, Inc.
STREET: 700 Capital Square, 400 Locust Street
STRET: Des Moines
STATE: IOWA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Diskette, 3.5 inch, 1.44 Mb storage
                                                                                SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/440,174A
FILING DATE: 12-MAY-1995
CLASSIFICATION: 530
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/079,512
FILING DATE: 18-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Yates, Michael E.; Sweeney, Patricia A.;
NAME: Roth, Michael J.; & Simon, Soma G.
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb st
COMPUTER: IBM Compatible
COMPUTER: SYSTEM: MS-DOS/Microsoft Windows
SOFRATING SYSTEM: MS-DOS/Microsoft Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/00062
                                                                                                                                                                                                                                                                                                                            0234R2D-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 234R2-PCT TELECOMMUNICATION: INFORMATION: TELEPHONE: (515) 248-4800
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application PC/TUS9500062 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                      NAME: BODTOWICZ, DODINA
REGISTRATION NUMBER: 32,196
REFRENCE/DOCKET NUMBER: 02341
TELECOMMUNICATION INFORMATION:
TELEPAX: (515) 248-4896
TELEPAX: (515) 334-6883
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25.00.
100.08; Fix
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INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KY: United States 50309
                                                                                                                                                                                                                                                                                                                                                                                                                                                         38 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: protein US-08-440-174A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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Matches
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Sequence 29973, Application US/09252991A

TITLE OF INVENTION:
TITLE APPLICATION NUMBER: US 60/074,788

PRIOR FILING DATE: 1998-02-18

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 29973

LENGIN:
TENTIOR TITLE OF INVENTION UNMER IN THE OF THE OF
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APPLICANT: Dumas Mine Edwards, J.B.
APPLICANT: Dumas Mine Edwards, J.B.
APPLICANT: Dumas Mine Edwards, J.B.
APPLICANT: Giordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.02.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR PILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                   25.0%; Score 5; DB 3; Length 61; 100.0%; Pred. No. 1.4e+02; rive 0; Mismatches 0; Indels
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1.4e+02;
0;
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OTHER INFORMATION: score 4
OTHER INFORMATION: seq LLQELGLCMCLLS/AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 4436, Application US/09513999C Patent No. 6783961
                                                                                              ; TYPE: PRT
; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-3836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5; Conservative
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NUMBER OF SEQ ID NOS: 5674
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ORGANISM: Homo sapiens
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Best Local Similarity
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                      11 RWLRR 15
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NAME/KEY: SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 47
US-09-513-999C-4436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-513-999C-4436
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                              SEQ ID NO 3836
LENGTH: 61
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FILE REPERENCE: GTC-007

CURRENT APPLICATION NUMBER: US 60/064,964

PRIOR FILING DATE: 1997-11-08

PRIOR FILING DATE: 1997-08-14
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                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz & Norris
STREET: One Liberty Place 46th Floor
CITY: Philadelphia
CITY: Pensylvania
COUNTRY: United States of America
ZIP: 19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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                                                                                                                                                                                               Sequence 1, Application PC/TUS9414074
GENERAL INFORMATION
APPLICANT: Maksymowych, Andrew B
APPLICANT: Hsu, Tin-Chen
APPLICANT: Jameson, Bradford A
APPLICANT: Jivack, Gerald
TITLE OF INVENTION: Biologically Active Compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER. READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURREMY APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASIPECATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/164,102
FILING DATE: 07-DEC-1993
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: DeLuca, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-134-001C-3836
; Sequence 3836, Application US/09134001C
; Patent No. 6380370
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 60 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5; Conservative
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MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
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Best Local Similarity
Matches 5; Conserv
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55 RYGWN 59
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GENERAL INFORMATION:
APPLICANT: Gary L. Breton et al.
APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO MORAXELLA CATARF
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REPERRICE: 2709.2005-001
CURRENT APPLICATION NUMBER: US/09/540,236
CURRENT FILING DATE: 2000-04-04
NUMBER OF SEQ ID NOS: 3840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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TITLE OF INVENTION: 207 Human Secreted Proteins
FILE REFERENCE: PZ007P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT PEPLICATION NUMBER: US/09/205, 258

CURRENT FILING DATE: 1998-12-04

EARLIER APPLICATION NUMBER: DCT/US98/11422

EARLIER APPLICATION NUMBER: 60/049, 885

EARLIER FILING DATE: 1997-06-06

EARLIER FILING DATE: 1997-06-06

EARLIER PILING DATE: 1997-06-06

EARLIER PILING DATE: 1997-06-06

EARLIER FILING DATE: 1997-06-06

EARLIER PELING DATE: 1997-06-06

EARLIER FILING DATE: 1997-06-06

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100.0%; Pred. No. 1.7
tive 0; Mismatches
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APPLICATION NUMBER: 60/048,971
FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,964
FILING DATE: 1997-06-06
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FILING DATE: 1997-06-06
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FILING DATE: 1997-06-06
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EARLIER APPLICATION NUMBER: 60/048,892
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EARLIER APPLICATION NUMBER: 60/048,915
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Patent No. 6525174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 1997-06-06
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Best Local Similarity
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APPLICANT: Young et
                                                                                                                                                                                                                                                                                                                                                                                           US-09-540-236-2141
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                  Patent No.
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Sequence 6143, Application US/09513999C

Patent No. 6783961

APPLICANT: Dumas Milne Edwards, J.B.

APPLICANT: Duclert, A.

APPLICANT: Giordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

Patent No. 6783961

Patent No. 6783961

CURRENT APPLICATION NUMBER: US/09/513,999C

PRIOR APPLICATION NUMBER: US 60/122,487

PRIOR FILING DATE: 1999-02-26

NUMBER OF SEQ ID NOS: 36681

SOFTWARE: Patent.pm

SEQ ID NO 6143
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APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: FILE REFERENCE: FILE REFERENCE: 1326-094
CURRENT APPLICATION NUMBER: US/09/270,767
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 59078
LENGTH: 79
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100.0%; Pred. No. 1.6e+02;
ive 0; Mismatches 0;
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100.0%; Pred. No. ...
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100.0%; Pred. No. 1.6
tive 0; Mismatches
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Patent No. 6703491
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US-09-540-236-2141
; Sequence 2141, Application US/09540236
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ORGANISM: Drosophila melanogaster
                                                                                                                  5; Conservative
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Best Local Similarity 100.
Matches 5; Conservative
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Matches 5; Conservative
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CORGANISM: Homo sapiens
US-09-513-999C-6143
                                                   Query Match
Best Local Similarity
                                                                                                                                                                        11 RWLRR 15
                                                                                                                                                                                                                              10 RWLRR 14
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US-09-252-991A-29973
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GENERAL INFORMATION:

APPLICANT: GETY 1. Ereton et al.

APPLICANT: GETY 1. Ereton et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO MORAXELLA CATAR

TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 2709-2005-001

CURRENT APPLICATION NUMBER: US/09/540,236

CURRENT FILING DATE: 2000-04-04

NUMBER OF SEQ ID NOS: 3840

SEQ ID NO 2144

LENGTH: 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Farb, David H
APPLICANT: Russek, Shelley
APPLICANT: Globs, Terrell
TITLE OF INVENTION: EFFECT OF STEROIDS ON NMDA RECEPTORS DEPENDS ON SUBUNIT
TITLE OF INVENTION: COMPOSITION
FILE REFERENCE: 0146-2026
CURRENT APPLICATION NUMBER: US/09/652,345
CURRENT APPLICATION NUMBER: US/09/652,345
PRIOR APPLICATION NUMBER: 00/151,802
PRIOR APPLICATION NUMBER: 09/151,802
PRIOR FILING DATE: 1999-08-31
PRIOR FILING DATE: 1999-08-31
PRIOR FILING DATE: 1999-08-20
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PALENTIN OF: 2.0
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100.0%; Pred. No. 1.9e+(
ative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-652-345-4; Sequence 4, Application US/09652345; Patent No. 6623933
  1998-04-20
PRIOR FILING DATE: 1998-04-2
NUMBER OF SEQ ID NOS: 107
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 25
LENGTH: 92
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, ORGANISM: M.catarrhalis
US-09-540-236-2144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                        TYPE: PRT; ORGANISM: Homo sapiens
US-09-673-809-25
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Best Local Similarity
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Best Local Similarity
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US-09-540-236-2144
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LOCATION: (15)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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Patent No. 6528261
GENERAL INFORMATION:
GENERAL INNOGENETICS N.V.
TITLE OF INVENTION: Method for typing of HLA alleles.
FILE REPERENCE: PCT99.86.HLA
CURRENT APPLICATION NUMBER: US/09/673,809
CURRENT FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 98870088.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EARLIER FILING DATE: 1997-06-06
EARLIER PILING DATE: 1997-06-06
EARLIER PILING DATE: 1997-06-06
EARLIER FILING DATE: 1997-06-06
EARLIER APPLICATION NUMBER: 60/048,878
EARLIER APPLICATION NUMBER: 60/070,923
EARLIER APPLICATION NUMBER: 60/092,921
EARLIER APPLICATION NUMBER: 60/092,921
EARLIER APPLICATION NUMBER: 60/092,921
EARLIER APPLICATION NUMBER: 60/094,657
EARLIER FILING DATE: 1998-07-15
EARLIER FILING DATE: 1998-07-30
                                                                                        R PILLING DATE: 1997-06-06
R APPLICATION NUMBER: 60/048,972
R FILING DATE: 1997-06-06
R PILING DATE: 1997-06-06
R FILING DATE: 1997-06-06
R APPLICATION NUMBER: 60/049,373
      FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/049,019
FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,970
                                                                                                                                                                                                                                                                    FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/049,374
FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,917
FILING DATE: 1997-06-06
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APPLICATION WUMBER: 60/048,949
ELING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,974
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APPLICATION WUMBER: 60/048,883
FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,897
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APPLICATION NUMBER: 60/048,898
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APPLICATION NUMBER: 60/048,962
FILING DATE: 1997-06-06
APPLICATION NUMBER: 60/048,963
FILING DATE: 1997-06-06
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APPLICATION NUMBER: 60/048,875
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SOFTWARE: PatentIn Ver. 2.0
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Best Local Similarity
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US-09-673-809-25
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LENGTH: 82
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         BARLIER P
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Sequence 37133, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: U5/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 52350
LENGTH: 143
                                                                                                                                                                                                                                                                                                                                                                      Gaps
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    REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET WINGBER: 230/045
TELECOMMUNICATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELERX: 67-3510
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
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12-09-270-767-52350
1. Sequence 52350, Application US/09270767
1. Patent No. 6703491
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ORGANISM: Drosophila melanogaster
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US-09-270-767-37133
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    Indels
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Patent No. 6310181
GENERAL INFORMATION:
APPLICANT: Kouhara, Haruhiko
APPLICANT: Spivak-Kroizman, Taly
APPLICANT: Schlessinger, Joseph
TITLE OF INVENTION: ADAPTOR PROTEIN FRS2 AND
TITLE OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                               APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SEQ ID NO 5186
LENGTH: 119
    0;
  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
CUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM COMPatible
COMPUTER: IBM COMPatible
COMPUTER: IBM COMPatible
COMPUTER: BEASEQ for Windows 2.0
SOFTWARE: FeatsERQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/980,523
FILING DATE: December 1, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCT/US97/21851
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8.09-621-976-5186
Sequence 5186, Application US/09621976
Patent No. 6639063
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APPLICANT: Dumas Milne Edwards, J.B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/US97/21
FILING DATE: December 1, 1997
APPLICATION NUMBER: 60/032,093
FILING DATE: December 3, 1996
ATTORNEY, AGENT INFORMATION:
NAME: Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 5; Conservative
5; Conservative
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
                                       6 WALGW 10
                                                                              11 WALGW 15
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Matches
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GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: ACID AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1999-07-27

NUMBER OF SEQ ID NOS: 33142

LENGTH: 158
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APPLICANT: Gordon G.
APPLICANT: Clark, Hilary
APPLICANT: Clark, Hilary
APPLICANT: Genetic, Kim
APPLICANT: Genetics Institute, Inc.
TILIB OF INVENTION: SECRETAD PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REPRENCE: GI 63300-11A
CURRENT APPLICATION NUMBER: US/09/311,021
CURRENT PILING DATE: 1999-05-13
NUMBER OF SEQ ID NOS: 268
SOFTWARE: Patentin Ver. 2.0
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Pred, No. 2.8e+02;
0; Mismatches 0; Indels
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                                                                                                                                                                    DB 4; Length 145; . 2.7e+02;
                                                                                                                                                                                                          0; Indels
                                                                                                                                                                    25.0%; Score 5; DB 4
100.0%; Pred. No. 2.7
tive 0; Mismatches
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; Patent No. 6706869
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100.0%; Pre
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                                                                                                        ORGANISM: Pseudomonas aeruginosa
1998-07-27
PRIOR FILING DATE: 1998-07-2
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 28008
LENGTH: 145
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Best Local Similarity 100.0
Loc 5; Conservative
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Best Local Similarity 100..
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Best Local Similarity 100.
Matches 5; Conservative
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US-09-252-991A-27239
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LENGTH: 149
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                                                                                    TYPE: PRT
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GENERAL INFORMATION:

APPLICANT: MAIC J. Rubenfield et al.

APPLICANT: MAIC J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS

FILLE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT APPLICATION NUMBER: US/09/27108
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Patent No. 6703491

GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster; FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767

CURRENT FILING DATE: 1999-03-17

SOFTWARE: PatentIN Vor. 2.0

SEQ ID NO 47237
                                                                                                                                                                                                            APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILLE REFERENCE: FALE Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 32020
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PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
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Patent No. 6551795
                                                                                                                                   US-09-270-767-32020

. Sequence 32020, Application US/09270767

. Patent No. 6703491

. GRMERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Drosophila melanogaster
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ORGANISM: Drosophila melanogaster
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        RYGWN 19
                                              RYGWN 53
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US-09-252-991A-28008
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APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: ARRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REPERBACE: 107196,136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR PRILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 32019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: ARRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NOS: 33142
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FAUCHT WOOD STATES

APPLICANT: MARC J. Rubenfield et al.

APPLICANT: MARC J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: ABENGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR FILING DATE: 1998-02-18

PRIOR FILING DATE: 1998-07-18

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142
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100.0%; Pred. No. 3.1
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                          ) ORGANISM: Pseudomonas aeruginosa US-09-252-991A-32019
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Best Local Similarity 100.
Matches 5; Conservative
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Best Local Similarity
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CRGANISM: Pseudor
US-09-252-991A-18579
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                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MAIC J. Rubenfield et al.
APPLICANT: MAIC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: ABENGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196_136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 20364
LENGTH: 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 30988, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION: WILLIAM APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS;
TITLE OF INVENTION: ARROGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: ARROGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT PELLING DATE: 1999-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR PELLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25.0%; Score 5; DB 4; Length 168; 100.0%; Pred. No. 3.1e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                               Sequence 20364, Application US/09252991A Patent No. 6551795
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Pseudomonas aeruginosa
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Matches 5; Conserv
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Best Local Similarity
                          RWLRR 15
                                                                    21 RWLRR 25
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US-09-252-991A-32019
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LENGTH: 168
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Patent No. 5451512

GENERAL INFORMATION:
APPLICANT: Apple, Raymond J.
APPLICANT: Bugawan, Teodorica L.
APPLICANT: Erlich, Henry A.
TITLE OF INVENTION: Methods and Reagents for HLA Class I A
TITLE OF INVENTION: Locus DNA Typing
NUMBER OF SEQUENCES: 173
                                                                                                    US-08-127-954-135

Sequence 135, Application US/08127954

Patent No. 5451512

GENERAL INFORMATION:
APPLICANT: Apple, Raymond J.
APPLICANT: Bugawan, Teodorica L.
APPLICANT: Exlich, Henry A.
TITLE OF INVENTION: Methods and Reagents for HLA Class I A
TITLE OF INVENTION: Locus DNA Typing
NUMBER OF SEQUENCES: 173
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25.0%; Score 5; DB 1; Length 182; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/127,954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffmann-La Roche Inc.
STREET: 340 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Hoffmann-La Roche Inc. STREET: 340 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas A.
REGISTRATION NUMBER: 35,321
REPERENCE/DOCKET NUMBER: 8873
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (510) 814-2977 INFORMATION FOR SEQ ID NO: 135:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: DNA (genomic) US-08-127-954-135
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amino acid
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Best Local Similarity 100.
Matches 5; Conservative
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              147 LRRYG 151
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CITY: Nutley
STATE: New Jer
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US-08-127-954-137
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Sequence 17721, Application US/09248796A
Sequence 17721, Application US/09248796A
Sequence 17721, Application US/09248796A
Sequence 17721, Application US/09248796A
TITLE OF INVENTION: WOLLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: WOLLEIC ACID AND THERAPEUTICS
FILE REPERENCE: 107196-132
FILE REPERENCE: 107196-132
PRIOR APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1998-02-12
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
LENGTH: 180
                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER: OF SEQ. ID NOS: 33142
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                                                                                                                                   25.0%; Score 5; DB 4; Length 173; 100.0%; Pred. No. 3.2e+02; ive 0; Mismatches 0; Indels
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                                        TYPE: PRT ORGANISM: Pseudomonas aeruginosa
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                                                                                                             Query Match
Best Local Similarity luv...
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US-09-248-796A-17721
                                                                                        US-09-252-991A-20339
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US-09-252-991A-24281
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LENGTH: 176
SEQ ID NO 20339
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                       LENGTH:
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COUNTRY:

13 LRRYG 17

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Sequence 139, Application US/08127954
Patent No. 5451512
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Apple, Raymond J.
APPLICANT: Apple, Raymond J.
APPLICANT: Enlich, Henry A.
TITLE OF INVENTION: Methods and Reagents for HLA Class I A
TITLE OF INVENTION: Locus DNA Typing
NUMBER OF SEQUENCES: 173
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffmann-La Roche Inc.
STREET: 340 Kingsland Street
                                                                                                                                                                                                          25.0%; Score 5; DB 1; Length 182; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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STATE: New Jersey
COUNTRY: U.S.A.
ZIP: 07110-1199
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BR PC computeR: BR PC computeR: B
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2974
TELEFAX: (510) 814-2977
INFORMATION FOR SEQ ID NO: 139:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              35,321
                                   TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
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CLASSIFICATION: 436
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas A.
REGISTRATION NUMBER: 35,2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 182 amino acids
TYPE: amino acid
                                                                                                                                                                  Query Match
Best Local Similarity 100.0
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STRANDEDNESS: single TOPOLOGY: linear
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                      12 WLRRY 16
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                                                                                                                           US-08-127-954-138
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US-08-127-954-139
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US-08-127-954-139
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US-08-127-954-138
Sequence 138, Application US/08127954
FATELL SASSISIZ
GENERAL INFORMATION:
APPLICANT: Apple, Raymond J.
APPLICANT: Bugawan, Teodorica L.
APPLICANT: Enlar, Henry A.
TITLE OF INVENTION: Methods and Reagents for HLA Class I A.
TITLE OF INVENTION: Locus DNA Typing
NUMBER OF SEQUENCES: 173
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25.0%; Score 5; DB 1; Length 182; 100.0%; Pred. No. 3.3e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                    SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/127,954
FILING DATE:
CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: PETRY DOUGHSR: 35,321
REFERENCE/DOCKET NUMBER: 35,321
REFERENCE/DOCKET NUMBER: 36,321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/127,954
FILING DATE:
CLASSIETCATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: PRETY, Douglas A.
REGISTRATION NUMBER: 35,321
REFERENCE/DOCKET NUMBER: 8873
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Hoffmann-Ia Roche Inc.
STREET: 340 Kingsland Street
CITY: Nutley
STATE: New Jersey
COUNTRY: U.S.A.
ZIP: 07110-1139
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATING SYSTEM:
OPERATING SYSTEM: PC-DOS/MS-DOS
             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATER: PATENT PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0.
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TELEFAX: (510) 814-2977
INFORMATION FOR SEQ ID NO: 138:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-127-954-137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 182 amino acids
TYPE: amino acid
STRANDEDNESS: single
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amino acid
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Best Local Similarity
Matches 5; Conserv
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TYPE: am
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Gaps

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Title: Perfect score:

Sequence:

OM protein

Run on:

Scoring table:

Searched:

Total number Word size :

Minimum DB s Maximum DB s

Database :

Score

Result No.  $\mathbf{v}$   $\mathbf{v}$ 

hypothetical prote probable ketoacyl MFC HLA-Azalpha-2 MHC H2-K transplan MHC K-bm6 transplan MHC class I H2-K-b bombyxin A-3 homol bombyxin A-1 homol hypothetical prote phosphotyrosine prepatch repair prote hypothetical prote hypothetical prote hypothetical prote hypothetical prote probable flavoprot hypothetical prote hypothetical prote hypothetical prote hypothetical prote hypothetical prote hypothetical prote ubiquinol-cytochro conserved hypothetical prote hypothetical prote class I histocompa marc class I antige MHC class I antige MHC class I antige MHC class I antige hypothetical prote hypothetical prote pathogenesis-relat hypothetical prote hypothetical prote MMC R121 protein -MMC class I histoc MMC cell surface a MMC cell surface g hypothetical prote hypothetical prote human leukocyte an Ig heavy chain V r hypothetical prote Na+/H+ antiporter hypothetical prote protein C48B4.4b [ trans-regulatory s trans-regulatory s finger protein HPF histocompatibility histocompatibility Staphylococcus two Staphylococcus two HLA-Cw7 - human (f transport protein conserved hypothet conserved hypothet ribosomal protein asparagine-tRNA li probable membrane-type I keratin 48k hypothetical prote conserved hypothet hypothetical prote hypothetical prote

## A;Residues: 1-142 <SAK> Matches g $\overset{>}{\circ}$ a ò C;Species: Marmota monax (woodchuck) C;Species: Marmota monax (woodchuck) C;Date: 02-ul-1996 #sequence\_revision 02-uul-1996 #text\_change 09-uul-2004 C;Accession: 148195 R;Yamazoe, M.; Nakai, S.; Ogasawara, N.; Yoshikawa, H. R;Yamazoe, M.; Nakai, S.; Ogasawara, N.; Yoshikawa, H. R;Title: Integration of woodchuck hepatitis virus (whv) DNA at two chromosomal sites (v-A;Title: Integration of woodchuck hepatitis virus (whv) DNA at two chromosomal sites (v-A;Reference number: 148190; MUID:91276235; PMID:2055466 A;Recession: 148192 A;Accession: preliminary; translated from GB/BMBL/DDBJ A;Residues: 1-123 <RES. A;Molecule type: DNA A;Residues: 1-123 <RES. A;Across-references: UNIPROT:090137; GB:M60766; NID:g191471; PIDN:AAA37108.1; PID:g191472 C;Superfamily: hepatitis B virus surface antigen C;Keywords: surface antigen hypothetical protein PA2661 [imported] - Pseudomonas aeruginosa (strain PAO1) C;Species: Pseudomonas aeruginosa C;Species: Pseudomonas aeruginosa C;Species: Dsep-2000 #sequence\_revision 15-Sep-2000 #text\_change 09-Jul-2004 C;Accession: C83311 R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Braman, S.; Yuan, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Brancer, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Brancer, G.S.; Olson, M.V. S.; Olson, M.V. S.; Olson, M.V. Stritle: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathon, A;Reference number: A82950; MuID:20437337; PMID:10984043 A;Accession: C83311 A;Accession: C83311 A;Accession: C83311 A;Accession: Preliminary A;Molecule type: DNA A;Residues: 1-297 <STO> A;Residues: 1-297 <STO> A;Residues: 1-297 <STO> A;Residues: 1-297 <STO> A;Residues: 1-207 <STO> A;Residues: 148552 orf US426 - infectious laryngotracheitis virus C;Specia: infectious laryngotracheitis virus C;Date: 17-Feb-1994 #sequence\_revision 17-Feb-1994 #text\_change 08-Oct-1999 C;Accession: 148552 C;Accession: 148552 R;Sakaguchi, M.; Urakwa, T.; Hirayama, Y.; Miki, N.; Yamamoto, M.; Hirai, K. Virus Genes 6, 365-378, 1992 A;Title: Sequence determination and genetic content of an 8.9-kb restriction fragment: A;Reference number: A48552; MUID:93118245; PMID:1282282 0 ö Gaps . 0 ; 0 Length 123; Length 297; 0; Indels Indels 0 DB 2; . 0.13; 30.0%; Score 6; DB 2; llarity 100.0%; Pred. No. 10; Conservative 0; Mismatches 40.0%; Score 8; DB 2 100.0%; Pred. No. 0.1 iive 0; Mismatches ALIGNMENTS surface antigen - woodchuck (fragment) Best Local Similarity 100. Matches 8; Conservative 8 LGWRWLRR 15 72 LGWRWLRR 79 Local Similarity SSWALG 56 4 SSWALG 9 Status: preliminary A; Molecule type: DNA A; Gene: PA2661 Query Match A; Accession: A; Status: pre Query Match RESULT 3 RESULT 1 q à d

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Cippecies: Bacillus subtlis:
Cippecies: Branch S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chor, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chor, S.; Bruschi, C.V.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E. Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Funa, S.; Hosono, S.; Hullo, M.F.; Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, K.; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueell, Y.; M.; Ogawar, K.; Ogiwara, A.; Odiega, B.; Bark, S.H.; Sarro, V.; Pohl, T.M.; Portetelle, Y.; M.; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seror A;Authors: Schleich, S.; Schroeter, R.; Torsato, V.; Popliyama, A;Authors: Yoshikawa, H.F.; Zumstein, B.; Yoshikawa, H.; Danchin, A.; Tosato, V.; Uchiyama, A;Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.; Tosato, V.; Uchiyama, A;Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.; Atata, K.; Yashancto, M.; A;Accession: C69780
A;Accession: C69780
A;Accession: C69780
A;Accession: C69780
A;Accession: C69780
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O
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C;Date: 04-Feb-2000 #sequence_revision 04-Feb-2000 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Ottenwaelder, B.; Obermaier, B.; Mewes, H.W.; Gassenhuber, J.; Wiemann, submitted to the Protein Sequence Database, January 2000
A;Reference number: Z23031
A;Accession: T46382
A,Cross-references: GB:M80595; NID:g291557; PIDN:AAB59898.1; PID:g291567
A,Note: sequence extracted from NCBI backbone (NCBIN:121622, NCBIP:121633)
                                                                                                                                                                            Gaps
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                                                                                                      Length 142;
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A,Cross-references: UNIPROT:Q86T03; EMBL:AL137727
A,Experimental source: adult testis; clone DKFZp434M0519
C,Genetics:
A,Note: DKFZp434M0519.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein DKFZp434M0519.1 - human (fragment)
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                                                                                                              DB 2;
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                                                                                                              30.0%; Score 6; DB 2;
100.0%; Pred. No. 12;
:ive 0; Mismatches
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Matches 6; Conserv
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Best Local Similarity
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Length 207;

DB 2;

Score 6;

30.0%;

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R,Norder, H.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 3141-3145, 1992
A,Fille: Molecular basis of hepatitis B virus serotype variations within the four major s
A,Reference number: PQ0453; MUID:93107848; PMID:1469353
A,Accession: PQ0599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C'Accession: UQ2119; PQ0600 - T. Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, J. Ronder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, J. Gen. Virol. 74, 1341-1348, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Small surface protein - hepatitis B virus
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Sate: 19-May-1994 #sequence_revision 19-May-1994 #text_change 09-Jul-2004
C;Accession: JG2225; S31798
B;Naumann, H.; Schaefer, S.; Yoshida, C.F.T.; Gaspar, A.M.C.; Repp, R.; Gerlich, W.H.
J. Gen. Virol. 74, 1627-1632, 1993
A;Tile: Identification of a new hepatitis B virus (HBV) genotype from Brazil that express A;Reference number: JG2225; MUID:93346970; PMID:8345355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            surface antigen - hepatitis B virus (subtype indet, strain Man)
C;Species: hepatitis B virus, HBV
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                         C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
F;3/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                  Length 226;
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                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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. 16;
                                                                                                                                                                                  A, Experimental source: subtype adw4q-, strain Rou
C, Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A.Gene: S
C.Superfamily: hepatitis B virus surface antigen
C.Keywords: surface antigen
                                                                                                                                                                                                                                                                                                                                                             30.0%; Score 6; DB 1
100.0%; Pred. No. 16;
live 0; Mismatches
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Pred. No.
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100.0%; Pre-
0; h
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                                                                                                                                A; Molecule type: DNA
A; Residues: 101-180 <NO3>
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A;Molecule type: DNA
A;Residues: 101-180 <NO2>
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Best Local Similarity
Matches 6; Conserv
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A; Residues: 1-226 <NOR>
A; Residues: 1-226 <NOR>
A; Cross-references: UNTROT:069609
A; Cross-references: Untract adw4q, strain CNTS-38
B; Norder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, J. Gen. Virol. 74, 1341-1348, 1993
A; Title: Genetic relatedness of hepatitis B viral strains of diverse geographical origin A; Reference number: UQ2044; MUID:93329382; PMID:8336122
                                                                                                                                                                                                                                                                                                                                                C;Accession: JQ1579; JQ2117
R;Norder, H.; Hammas, B.; Loefdahl, S.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 1201-1208, 1992
A;Title: Comparison of the amino acid sequences of nine different serotypes of hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A.Cross-references: UNIPROT:069612; GB:X75661; NID:9416088; PIDN:CAA53348.1; PID:9416089
A;Experimental source: subtype adw4q-, strain CNTS-36
B;Norder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, J. G., J. 1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains of diverse geographical origin A;Reference number: UQ2044; MUD:93329382; PMID:8336122
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C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 09-Jul-2004
C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: JG1809, JQ2118; PQ0559
R;Norder, H.; Hammas, B.; Locfdahl, S.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 1201-1208, 1992
A;Title: Comparison of the amino acid sequences of nine different serotypes of hepatitis
A;Reference number: JQ1570; MUID:92268879; PMID:1588323
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C;Species: hepatitis B virus, HBV
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C;Species: hepatitis B virus, HBV
A;Note: host Homo sapiens (man)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
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                               Indels
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A,Molecule type: DNA
A,Residues: 1-226 <NO2>
A,Experimental source: subtype adw4q-, strain Car C,Genetics:
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A;Residues: 1-226 <NO2>
A;Experimental source: subtype adw4q-, strain Rou
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100.0%; Pred. No. 16;
Live 0; Mismatches
     Pred. No. 15;
                            Mismatches
                          0;
100.0%;
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Best Local Similarity 100...
                       Conservative
Best Local Similarity
Matches 6; Conserv
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Accession: JQ2118
                                                                         4 SSWALG 9
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R;Norder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, J. Gen. Virol. 74, 1341-1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains of diverse geographical origin A;Reference number: JQ2044; MUID:93329382; PMID:8336122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;CCCSS.referecs: UNIPROT:Q9B9A6; UNIPROT:Q8UMZ6; UNIPROT:Q9XXC1; UNIPROT:Q99HS3; UNIPPROT:Q91NX1; UNIPROT:Q91L64; UNIPROT:Q91LK5; UNIPROT:Q91LK64; UNIPROT:Q91LK65; UNIPROT:Q91LK64; UNIPROT:Q91LK65; UNIPROT:Q99HS8; UNIPROT:Q99HS8; UNIPROT:Q99HS8; UNIPROT:Q99HS9; UNIPROT:Q90HS9; UNIPROT
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C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
C;Accession: D83210
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Bradman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim, J. Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
A;Ritler Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic patho A;Reference number: A82950; MUID:20437337; PMID:10984043
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A;Experimental source: strain PAO1
C;Genetics:
A;Gene: PA3483
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C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Dates: 19-May-1994 #sequence_revision 19-May-1994 #text_change 09-Jul-2004
C;Dates: 19-May-1994 #sequence_revision 19-May-1994 #text_change 09-Jul-2004
C;Accession: JQ2226; S31795
R;Naumann, H.; Schaefer, S.; Yoshida, C.F.T.; Gaspar, A.M.C.; Repp, R.; Gerlich, W.H.
J Gen. Virol. 74, 1627-1632, 1993
A;Title: Identification of a new hepatitis B virus (HBV) genotype from Brazil that expre
A;Reference number: JQ2225; MUID:93346970; PMID:8345355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein PA3483 [imported] - Pseudomonas aeruginosa (strain PAO1)
                                                                                                                                                           surface antigen - hepatitis B virus (subtype adw4q-, strain 273/85)
C,Species: hepatitis B virus, HBV
C,Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
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C;Superfamily: hepatitis B virus surface antigen
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100.0%; Pred. No. 16;
Live 0; Mismatches
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100.0%; Pred. No. 19;
ive 0; Mismatches
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Best Local Similarity
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A; Residues: 1-226 < NOR>
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Suffaces thepatitis B virus, HBV
C; Deciss. hepatitis B virus, HBV
C; Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
C; Den. Virol. 74, 1341-1348, 1993
A; Center Genetic relatedness of hepatitis B viral strains of diverse geographical origin
A; Reference number: UQ2044; MUID:93329382; PMID:8336122
A; Contents: genogroup F
A; Accession: UQ2044
A; Residues: 1-226 *NOR
A; Residues: 1-226 *NOR
A; Residues: 1-226 *NOR
A; Residues: UNIPROT:Q99BA3; UNIPROT:Q99H83; UNIPROT:Q99H83; UNIPROT:Q99H83; UNIPROT:Q99H83; UNIPROT:Q99H83; UNIPROT:Q99H83; UNIPROT:Q99H89; UNIPROT:Q
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C;Species: hepatitis B virus, HBV
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
C;Cocession: J02021
R;Norder, H; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, G. Gen. Virol. 74, 1341-1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains of diverse geographical origin A;Reference number: J02044; MuID:9332982; PMID:8336122
A;Contents: genogroup F
A;Coctesion: J02021
A;Molecule type: DNA
A;Residues: 1-226 <NOR>
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A;Cross-references: UNIPROT:Q67870; EMBL:X69798; NID:g59422; PIDN:CAA49457.1; PID:g59428
C;Superfamily: hepatitis B virus surface antigen
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100.0%; Pred. No. 16;
iive 0; Mismatches
                                                                                                                                                                                           30.0%; Score 6; DB 2;
100.0%; Pred. No. 16;
iive 0; Mismatches
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C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen
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Similarity 100.0%; Pred. No. 16;
6; Conservative 0; Mismatches
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6, Conserve
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Tue Oct 26 08:02:17 2004

A;Accession: JQ2226

30.0%; Score 6; DB 2; 100.0%; Pred. No. 19; iive 0; Mismatches

Query Match Best Local Similarity 100. Matches 6; Conservative

209 SSWALG 214

4 SSWALG 9

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A;Residues: 1751 - CRI>
A;Cross-references: UNIPROT:P33586; GB:M93058; NID:g153396; PIDN:AAA72110.1; PID:g388262
C;Superfamily: type II thioesterase with ferredoxin-like domain; oleoy1-[acy1-carrier-prof: F;129-339/Domain: oleoy1-[acy1-carrier-prof: F;129-339/Domain: oleoy1-[acy1-carrier-prof: F;201/Active site: Ser #status predicted
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A;Nolecule type: DNDA
A;Residues: 1-379 <POL>
A;Ccoss-references: UNIPROT:Q56773; EMBL:U22511; NID:g1172090; PIDN:AAA86375.1; PID:g7333
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NiAltermate names: prest (LHRs) protein

C; Species: hepatitis B virus, HBV

C; Species: Jany-1994 #sequence_revision 19-May-1994 #text_change 09-Jul-2004

C; Accession: J02220; S11796

R; Naumann, H.; Schaefer, S.; Yoshida, C.F.T.; Gaspar, A.M.C.; Repp, R.; Gerlich, W.H.

J. Gen. Virol: 74, 1627-1632, 1932

A; Title: Identification of a new hepatitis B virus (HBV) genotype from Brazil that express

A; Reference number: J02225; MUID: 93346970; PMID: 83345355
                                                                                                                                                                                                                                                                                                                C,Accession: UN0530
R;Criado, L.M.; Martin, J.F.; Gil, J.A.
Gene 126, 135-139, 1993
A;Title: The pab gene of Streptomyces griseus, encoding p-aminobenzoic acid synthase, is
A;Reference number: JN0530; MUID:93231527; PMID:8472954
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                                                                                                                                                                                                           NiAlternate names: protein X
C;Species: Streptomyces griseus
C;Date: 31-Dec.1993 #sequence_revision 31-Dec-1993 #text_change 12-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GumG protein - Xanthomonas campestris
C;Species: Xanthomonas campestris
C;Date: 24-Aug-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S67856
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30.0%; Score 6; DB 2
Best Local Similarity 100.0%; Pred. No. 23;
Matches 6; Conservative 0; Mismatches
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submitted to the EMBL Data Library, March 1995
A.Reference number: 867856
A.Accession: 867856
                                                                                                                                                                                     thioesterase homolog - Streptomyces griseus
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C;Superfamily: O-acetyl transferase
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          23
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A, Molecule type: DNA
A, Residues: 1-400 < NAU>
       18 WRWLRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: DNA
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N/Contains: major surface antigen
C;Specials woodchuck hepatitis virus
C;Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 09-Jul-2004
C;Accession: B29498
R;Etiemble, J: Mercoey, T: Trepo, C:; Tiollais, P:; Buendia, M.A.
Gene 50, 207-214, 1986
A;Title: Nucleotide sequence of the woodchuck hepatitis virus surface antigen mRNAs and A;Reference number: A91568; MUID:87219879; PMID:3582979
A;Rocession: B29498
A;Molecule type: mRNA
A;Residues: 1-282 <ETI>A;Cross-references: UNIPROT:P11293; GB:M15954; NID:g893289; PIDN:AAA69574.1; PID:g336156
C;Genetics:
A;Cross-references: UNIPROT:P1293; GB:M15954; NID:g893289; PIDN:AAA69574.1; PID:g336156
C;Genetics: A;Genetics: B virus surface antigen
C;Keywords: glycoprotein; surface antigen
A;Molecule type: DNA
A;Residues: 1-281 <NAU>
A;Cross-references: UNIPROT:Q67869; EMBL:X69798; NID:g59422; PIDN:CAA49454.1; PID:g59425
C;Superfamily: hepatitis B virus surface antigen
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Lory, S.; Olson, M.V.
Altile: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic patholy. Riference number: Ag2950; MUID:20437337; PMID:10984043
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A;Molecule type: DNA
A;Residues: 1-331 <STO>
A;Cross-references: UNIPROT:0911W1; GB:AE004642; GB:AE004091; NID:g9948163; PIDN:AAG0554
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C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
C;Accession: A83377
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fl. 66/Domain: slyana sequence #status predicted <81G>
F:61-282/Product: major surface antigen (gene S) #status predicted <MSA>
F:3/Binding site: carbohydrate (Asn) (covalent) #status predicted
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DB 1;

30.0%; Score 6; DB 1; 100.0%; Pred. No. 19; tive 0; Mismatches

6; Conservative

Matches

Query Match Best Local Similarity

210 SSWALG 215

RESULT 16

4 SSWALG 9

à g DB 2;

22;

30.0%; Score 6; DB 2 100.0%; Pred. No. 22; tive 0; Mismatches

6; Conservative

10 WRWLRR 15

ò

Query Match Best Local Similarity Matches 6; Conserv

A,Experimental source: strain PAOI C,Genetics:

A; Gene: PA2154

A; Accession: A83377

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large surface antigen - ground squirrel hepatitis virus
N.Contains: major surface antigen; middle surface antigen
C;Species: ground squirrel hepatitis virus
C;Date: 25-Feb-1985 #sequence_revision 25-Feb-1985 #text_change 09-Jul-2004
C;Accession: A03709
R;Seeger, C; Ganem, D.; Varmus, H.E.
J. Virol. 51, 367-375, 1984
A;Itile: Nucleotide sequence of an infectious molecularly cloned genome of ground squirr
A;Reference number: A93000; MUID:84267998; PMID:6086950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      large surface antigen - woodchuck hepatitis virus (clone 59)
N;Contains: major surface antigen; middle surface antigen
C;Species: woodchuck hepatitis virus
C;Species: woodchuck hepatitis virus
C;Date: 30-Jun-1899 #sequence_revision 30-Jun-1989 #text_change 09-Jul-2004
C;Accession: H29969
R;Cohen, J.I.; Miller, R.H.; Rosenblum, B.; Denniston, K.; Gerin, J.L.; Purcell, R.H.
Virology 162, 12-20, 1989
A;Title: Sequence comparison of woodchuck hepatitis virus replicative forms shows conser
A;Reference number: A94368; MUID:88101359; PMID:3336938
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A,Molecule type: DNA
A,Residues: 1-431 <COHS
A,Cross-references: UNIPROT:P12910; GB:M19183; NID:g336141; PIDN:AAA46762.1; PID:g336142
C,Genetics:
A,Gene: pre-S1/pre-S2/S
C;Superfamily: hepatitis B virus surface antigen
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N;Contains: major surface antigen; middle surface antigen
C;Species: woodchuck hepatitis virus
C;Date: 30-Jun-1989 #sequence_revision 30-Jun-1989 #text_change 09-Jul-2004
C;Accession: D29969
R;Cohen, J.1.; Miller, R.H.; Rosenblum, B.; Denniston, K.; Gerin, J.L.; Purcell, R.H.
Virology 162, 12-20, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Gene: pre-S1/pre-S2/S
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
F;147-428/Product: middle surface antigen
F;207-428/Product: major surface antigen (gene S) #status predicted <MSA>
F;149/Binding site: carbohydrate (Asn) (covalent) #status predicted
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100.0%; Pred. No. 27;
ive 0; Mismatches
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100.0%; Pred. No. 27;
ative 0; Mismatches
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A,Residues: 1-428 <SES>
A,Cross-references: UNIPROT:P03144; GB:K02715
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Best Local Similarity
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Nothains: major surface antigen; middle surface antigen
C;Species: woodchuck hepatitis virus
C;Species: woodchuck hepatitis virus
C;Date: 14-Nov-1981 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A03707
R;Galibert, F: Chen, T.N.; Mandart, E.
J. Virol. 41, 51-65, 1982
A;Title: Nucleotide sequence of a cloned woodchuck hepatitis virus genome: comparison wing reference number: A92986; MUID: 82216969; PMID: 7086958
A;Accession: A03707
A;Molcule type: DNA
A;Residues: 1-426 cGL>
A;Cross-references: UNIPROT: P03143; GB: J02442; NID: G336126; PIDN: AAA46760.1; PID: G336128
C;Genetics:
A;Genetics:
A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics: A;Genetics:
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A;Residues: 1-411 <STO>
A;Residues: 1-411 <STO>
A;Residues: UNIPROT:Q9A3L1; GB:AE005673; NID:g13424868; PIDN:AAK25154.1; GSPDB:G
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            F87644

F87644

transcription regulator, TetR family [imported] - Caulobacter crescentus

transcription regulator, TetR family [imported] - Caulobacter crescentus

C;Species: Caulobacter crescentus

C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change 09-Jul-2004

C;Accession: F87644

R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, J.

B; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolon

H, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M.

Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001

A;Title: Complete Genome Sequence of Caulobacter crescentus.

A;Accession: F87644

A;Accession: F87644

A;Status: preliminary
A,Cross-references: UNIPROT:Q05496; EMBL:X69798; NID:g59422; PIDN:CAA49455.1; PID:g59426
C;Superfamily: hepatitis B virus surface antigen
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                                                                                                          30.0%; Score 6; DB 2
100.0%; Pred. No. 25;
tive 0; Mismatches
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nes 6; Conservative
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A)Cross_references: UNIPROT:P46882; EMBL:L38858; NID:g619754; PIDN:AAA98490.1; PID:g619754
A)Accession: S5267
A)Andlecule type: protein
B) A;Residues: 165-175 < SCW>
B;Schilling, B.; Lerch, K.
Biochim: Biophys. Acta 1243, 529-537, 1995
B) A;Title: Amine oxidases from Aspergillus niger: identification of a novel flavin-depender
B);Reference number: S60356; MUID:95244610; PMID:7727530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CjAccession: C84356

R;Ng, W.V.; Kennedy, S.P.; Mahairas, G.G.; Berquist, B.; Pan, M.; Shukla, H.D.; Lasky, S.; Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, D.W.; Maddocks, D.G.; Jablor Jung, K.H.; Alam, M.; Freitas, T.

Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000

A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.; Ebhardt, H.; Lowe, T.M.; Lishile: Genome sequence of Halobacterium species NRC-1.

A;Reference number: A84160; MUID:20504483; PMID:11016950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amine oxidase (flavin-containing) (EC 1.4.3.4) N - Aspergillus niger NyAlternate names: monoamine oxidase N C;Species Aspergillus niger C;Species Aspergillus niger C;Date: 14-Oct-1995 #sequence revision 12-Apr-1996 #text_change 09-Jul-2004 C;Accession: S55273; S55267; $60356 K;Schilling, B.; Lerch, K. Ms. Can. Genet. 247, 430-438, 1995 A;Title: Cloning, sequencing and heterologous expression of the monoamine oxidase gene A;Reference number: S55267; MUID:95287865; PMID:7770050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,Accession: S60356
A,Status: nucleic acid sequence not shown; not compared with conceptual translation
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C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
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F;41-69/Region: beta-alpha-beta FAD nucleotide-binding fold
F;493-495/Region: peroxisome/glyoxysome location signal #status atypical
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                                                                                                                                                                                                                                      Length 431;
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100.0%; Pred. No. 30;
tive 0; Mismatches
                  C;Superfamily: hepatitis B virus surface antigen
                                                                                                                                                                                                                           30.0%; Score 6; DB 1
100.0%; Pred. No. 27;
ative 0; Mismatches
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Best Local Similarity luv...
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459 WALGWR 464
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A; Residues: 1-495 <SCH>
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AsMolecule type: DNA
AsResidues: 1-571 <STO>
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C,Date: 30-Jun-1987 #sequence_revision 30-Jun-1987 #text_change 09-Jul-2004
C,Date: 30-Jun-1987 #sequence_revision 30-Jun-1987 #text_change 09-Jul-2004
C,Date: 30-Jun-1987 #sequence_revision 30-Jun-1987 #text_change 09-Jul-2004
C,Accession: A03708
A,Title: Nucleotide sequence of a cloned woodchuck hepatitis virus genome: evolutional r
A;Reference number: A93015; MUID:86062931; PMID:3855246
A;Reference number: A93015; MUID:86062931; PMID:3355346
A;Residues: 1-431 a and a and
A;Title: Sequence comparison of woodchuck hepatitis virus replicative forms shows consent A; Macession: D29969
A;Recession: D29969
A;Molecule type: DNA
A;Residues: 1-431 <COH>
A;COSS-references: UNIPROT:P12909; GB:M18752; NID:g336136; PIDN:AAA46766.1; PID:g336137
C;Genetics: A;Gene: pre-S1/pre-S2/S
C;Guperfamily: hepatitis B virus surface antigen
C;Keywords: g1/coprotein; surface antigen
C;Keywords: g1/coprotein; surface antigen (gene pre-S2/S) #status predicted <DSA>
F;20-431/Product: middle surface antigen (gene S) #status predicted <MSA>
F;210-431/Product: major surface antigen (gene S) #status predicted
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Ny Contains: major surface antigen, middle surface antigen
Cispecies: woodchuck hepatitis virus
Cispecies: woodchuck hepatitis virus
Cipate: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 09-Jul-2004
CiAccession: B32397
Cote, P.J. Hornbuckle, W.E.; Tennant, B.C.; Gerin, J.L.; Purcell, R.H.;
Proc. Natl. Acad. Sci. U.S.A. 86, 1846-1849, 1899
A;Title: Complete nucleotide sequence of a molecular clone of woodchuck hepatitis virus
A;Accession: B32397
A;Accession: B32397
A;Molecule type: DNA
A;Residues: 1-431 cdIR>
C;Genetics:
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100.0%; Pred. No.
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SAVLW8
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hypothetical protein P1.11659_3 - human
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A.Reference number: S74322; MUID:97061201; PMID:8905231
A.Accession: S76730
A.Status: preliminary
A.Molecule type: DNA
A.Molecule type: DNA
A.Residues: 1-636 < KAN>
A.Residues: 1-636 < KAN>
A.Residues: 1-636 < KAN>
A.Residues: 1-636 < KAN>
A.Residues: Leadences: UNIPROT:P74536; EMBL:D90916; GB:AB001339; NID:g1653715; PIDN:BAA1864
A.Rote: the nucleotide sequence was submitted to the EMBL Data Library, June 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypothetical protein - Synechocystis sp. (strain PCC 6803)

hypothetical protein - Synechocystis sp.

(Species: Synechocystis sp.

A;Varie: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004

C;Accession: S76730

C;Accession: S76730

C;Accession: S76730

C;Accession: J860

C;Accession: J87030

C;Accession:
A;Cross-references: UNIPROT:Q9HNKS; GB:AE004437; NID:g10581482; PIDN:AAG20215.1; GSPDB:G
C;Genetics:
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C:Species: Caenorhabditis elegans
C:Species: Caenorhabditis elegans
C:Species: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
C;Accession: T34232
R;Minx, P.
submitted to the EMBL Data Library, November 1995
A;Description: The sequence of C. elegans cosmid F20B6.
A;Accession: T34232
A;Access
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100.0%; Pred. No. 33;
tive 0; Mismatches
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Best Local Similarity
Matches 6; Conserv
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RESULT 31 T02245

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Cipecies: How sapiens (man)
Cipecies: How sapiens (man)
Cipecies: How sapiens (man)
Cipecies: Mar-1999 #sequence_revision 05-Mar-1999 #text_change 05-Nov-1999
Cipacession: 702245
Rilamerdin, J.E.; McCready, P.M.; Skowronski, E.; Adamson, A.W.; Burkhart-Schultz, K.; Gristensen, M.; Georgescu, A.; Avila, J.; Liu, S.; Attix, C.; Andreise, T.; Trankheir submitted to the EMBL Data Library, March 1998
A;Authors: Duarte, S.; Lucas, S.; Brucs, R.; Thomas, P.; Quan, G.; Kronmiller, B.; Arell A;Description: Sequence analysis of a human Pl clone containing the XRCC9 DNA repair gen A;Accession: T02245
A;Accession: T02245
A;Accession: MA
A;Accession: MA
A;Molecule type: DNA
A;Molecule type: DNA
A;Molecule type: EMBL;AC004472; NID:g2984582; PIDN:AAC07985.1; PID:g2984587
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A;Status: preliminary
A;Nolecule type: DNA
A;Cross-references: GB:Chr_III; PIDN:CAA82384.1; PID:g3875025; GSPDB:GN00021; CESP:C48B4
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A;Introns: 89/3; 152/1; 200/1; 241/2; 294/3; 354/3; 435/1; 733/1; 802/1; 871/1
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C,Species: Caenorhabditis elegans
C,Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 16-Aug-2004
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R,Kershaw, J.
R,Kershaw, J.
R,Kershaw, J.
R,Kershaw, J.
R,Reference number: S60124
A,Recession: S60124
A,Rocession: S60124
A,Molecule type: DNA
A,Molecule type: DNA
A,Korss-references: EMBL: Z29117; NID: 9439247; PID: 91066912
C;Genetics:
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100.0%; Pred. No. 46;
cive 0; Mismatches
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100.0%; Pred. No. 78;
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C,Superfamily: ATP-binding cassette homology
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position: III

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C;Species: Bacillus halodurans
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004
C;Accession: D83827
R;Takani, H.; Nakasone, K.; Takaki, Y.; Maeno, G.; Sasaki, R.; Masui, N.; Fuji, F.; Hiran Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and cancersion: D83827
A;Reference number: A88650; MUID:20512582; PMID:11058132
A;Accession: D83827
A;Actus: preliminary
A;Molecule type: DNA
A;Residues: 1-65 <STO>
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A;Experimental source: strain C-125
C;Genetics:
A;Gene: nahS
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C;Species: Aeropyrum pernix
C;Species: Aeropyrum pernix
C;Species: D-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
C;Accesion: A72733
R;Kawarabayasi, Y.; Hino, Y.; Horikawa, H.; Yamazaki, S.; Haikawa, Y.; Jin-no, K.; Takaha Awa, H.; Takamiya, M.; Masuda, S.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; KN DNA Res. 6, 83-101, 1999
A;Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyra A;Accession: A7273
A;Accession: A7273
A;Status: preliminary
A;Nolecule type: DNA
A;Residues: 1-75 <KAW>
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N;Alternate names: Lepb1170_C1_189 protein
C;Species: Mycobacterium leprae
C;Date: 19-Mar.1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: S72700
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100.0%; Pred. No. 90;
iive 0; Mismatches
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100.0%; Pred. No. 80;
iive 0; Mismatches
         25.0%; Score 5; DB 1100.0%; Pred. No. 76; ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 36
D83827
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                           A; interposes: 47/1; 112/3; 161/2; 220/2; 543/3; 574/3; 903/2; 1056/1; 1115/3; 1178/3; 1265/1; Superfamily: ATP-binding cassette homology
(Superfamily: ATP-binding cassette homology cassettes homology ca
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Theory chain V region (clone NP-7-5) - mouse (fragment)
C.Species: Mus musculus (house mouse)
C.Species: Mus musculus (house mouse)
C.Accession: PH1692
B;MCHeyzer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
B;MCHeyzer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
B;MCHeyzer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
A;McHeyzer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
A;McHeyzer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
A;Reference number: PH1675, MUID:93301607; PMID:8315385
A;Accession: PH1692
A;Accession: PH1692
A;Accession: PH1692
A;Accession: PH1692
A;Accession: Max
A;Residues: 1-23 ance;
A;Accession: Max
A;Residues: 1-23 ance;
A;Accession: McHibar (he codon AGT for residue 8 as Arg, GCT for residue 20 accession and accession and accession accession
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Date: 13-Jun-1983 #sequence revision 13-Jun-1983 #text_change 09-Jul-2004
Accession: C43011; H43016; Ā04393
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A,Start codon: GTG
C,Superfamily: phage lambda hypothetical 7.4K protein (nin region)
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At Reference number: A94614
At Accession: C43011
At Residues: 1-60 cDAN
At Residues: 1-60 cDAN
At Residues: 1-60 cDAN
At Cross-references: UNIPROT: P03768
B; Sanger F: Coulson, A.R.; Hong, G.F.; Hill, D.F.; Petersen, G.J. Mol. Biol. 162, 729-773, 1982
A Title: Nucleotide sequence of bacteriophage lambda DNA.
At Reference number: A92891; MUID: 83189071; PMID: 6221115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30.0%; Score 6; DB 2; Length 1767; 100.0%; Pred. No. 78; 0; Indels iive 0; Mismatches 0; Indels
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100.0%; Pred. No. 37;
iive 0; Mismatches
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Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity
Matches 5; Conserv
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C;Species: Mus musculus (house mouse)
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 09-Jul-2004
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 09-Jul-2004
C;Accession: 157611
R;Geliebter, J; Zeff, R.A.; Schulze, D.H.; Pease, L.R.; Weiss, E.H.; Mellor, A.L.; Flav Mol. Cell. Biol. 6, 645-652, 1986
A;Title: Interaction between K-b and Qg gene sequences generates the K-bm6 mutation.
A;Reference number: 157611; MUID:87064350; PMID:3023861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A, Cross-references: UNIPROT. 095555, GB.M12935; NID: 9199595; PIDN: AAA39677.1; PID: 9554238
C, Superfamily: class I histocompatibility antigen; immunoglobulin homology
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C;Species: Mus musculus (house mouse)
C;Date: 02-Uul-1996 #sequence_revision 02-Uul-1996 #text_change 09-Uul-2004
C;Accession: 159068
R;Geliebter, U; Zeff, R.A.; Melvold, R.W.; Nathenson, S.G.
R;Geliebter, U; A.3 Sci. U;S.A. 83, 3371-3375, 1986
Proc. Natl. Acad. Sci. U;S.A. 83, 3371-3375, 1986
A;Titler: Micotic recombination in germ cells generated two major histocompatibility comp A;Reference number: 159068; MUID:86205899; PMID:3458188
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A;Cross-references: UNIPROT:Q31189; GB:M13200; NID:g199535; PIDN:AAA39649.1; PID:g554230
C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
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bombyxin A-3 homolog - ailanthus silkmoth
bombyxin A-3 homolog - ailanthus silkmoth)
C;Species: Samia cynthia (ailanthus silkmoth)
C;Date: 112-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
R;Iwami, M:
R;Iwami, M:
A;Reference number: JQ0902
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A,Molecule type: DNA
A,Residues: 1-98 - SIWA
A,Cross-references: UNIPROT: P33720
A,Note: the authors translated the codon ATA for residue 32 as Thr
A,Note: the source is designated as Samia cynthia ricini
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Mo. 16+02;
O; Indels
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                                                                                                                                                                                                157611
MHC K-bm6 transplantation antigen - mouse (fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-91 <RES>
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Best Local Similarity 100.0%; Pred. No. 16+
Matches 5; Conservative 0; Mismatches
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               12 WLRRY 16
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R,Sire, J.; Chimini, G.; Boretto, J.; Toubert, A.; Kahn-Perles, B.; Layet, C.; Sodoyer, J. Tumnol. 156009
R,Sire, J.; Chimini, G.; Boretto, J.; Toubert, A.; Kahn-Perles, B.; Layet, C.; Sodoyer, J.; Time: Hybrid genes between HLA-A2 and HLA-A3 constructed by in vivo recombination all A, Reference number: 156009; MUD: 88170828; PMID: 2450922
A, Accession: 156009
A, Steatus: preliminary; translated from GB/EMBL/DDBJ
A, Molecule type: DNA
A, Residues: 1-89 cRBs
A, Cross-references: UNIPROT: Q30176; GB: M20139; NID: g188493; PIDN: AAS59838.1; PID: g553587
C, Genetics: GDB: HLA-A
A, Cross-references: GDB: 119310; OMIM: 142800
A, Map position: 6p21.3-6p21.3
C, Superfamily: class I histocompatibility antigen; immunoglobulin homology
R;Smith, D.R.; Robison, K.
submitted to the EMBL Data Library, November 1993
A;Description: Mycobacterium leprae cosmid B1170.
A;Reference number: $72693
A;Accession: $72700
A;Status: preliminary
A;Status: preliminary
A;Residues: 1-84 <SMI>
A;Residues: 1-84 <SMI>
A;Cross-references: UNIPROT:Q49621; EMBL:U00010; NID:g466780; PIDN:AAA17064.1; PID:g4667
A;Genetics:
A;Genetics:
A;Genetics:
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C;Species: Mus musculus (house mouse)
C;Species: Oz-Uul-1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
C;Accession: 158989
R;Schulze, D.H.; Pease, L.R.; Geier, S.S.; Reyes, A.A.; Sarmiento, L.A.; Wallace, R.B.;
Proc. Natl. Acad. Sci. U.S.A. 80, 2007-2011, 1983
A;Title: comparison of the cloned h-2kbml variant gene with the h-2kb gene shows a clust A;Reference number: 158989; MUID:83169810; PMID:6300887
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A,Status: preliminary; translated from GB/EMBL/DDBJ
A,Molecule type: DNA
A,Molecule type: DNA
A,Ressidues: 1-91 «RES
A,Cross-references: UNIPROT:031190; GB:J00401; NID:g199540; PIDN:AAA39651.1; PID:g554232
C,Superfamily: class I histocompatibility antigen; immunoglobulin homology
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MHC HLA-A2-alpha-2 protein - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
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25.0%; Score 5; DB 2; Length 89;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 5; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 98;
iive 0; Mismatches
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Best Local Similarity 100.
Matches 5; Conservative
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C, Genetics:

RESULT 44

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Gispecies: Deinococcus radiodurans
Cispecies: Deinococcus radiodurans
Cispecies: Deinococcus radiodurans
Ciscession: H75591
Rsequence_revision 03-Dec-1999 #text_change 09-Jul-2004
CiAccession: H75591
Fighite, 0.; Eisen, J.A.; Heidelberg, J.F.; Hickey, E.K.; Peterson, J.D.; Dodson, R.J.; F.White, O.; Eisen, J.A.; Heidelberg, J.F.; Hickey, E.K.; Peterson, J.D.; Dodson, R.J.; F.White, S.Sauth, H.O.; Venter, J.C.; Fraser, C.M.
Soience 286, 1571-1577, 1999
A;Title: Genome sequence of the radioresistant bacterium Deinococcus radiodurans R1.
A;Reference number: A75250; MUID:20036896; PMID:10567266
probable flavoprotein [imported] - Clostridium acetobutylicum
C.Species: Clostridium acetobutylicum
C.Species: Clostridium acetobutylicum
C.Species: Clostridium acetobutylicum
C.Accession: G.P7084
R.Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee,
J. Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001
A.Fille: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium Clos
A.Reference number: A96900; MUID:21359325; PMID:21359325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-116 <WHI>
A;Cross-references: UNIPROT:Q9RYF3; GB:AE001863; GB:AE001825; NID:g6460670; PIDN:AAF1250;
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:Q971Y8; GB:AE001437; PIDN:AAK79466.1; PID:g15024445; GSPDB:GR
A;Experimental source: Clostridium acetobutylicum ATCC824
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A; Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyra A; Reference number: A72450; MUID:99310339; PMID:10382966
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C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
C;Accession: H72489
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C;Superfamily: Deinococcus radiodurans hypothetical protein DRA0365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein DRA0365 - Deinococcus radiodurans (strain R1)
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Best Local Similarity 100.
Matches 5; Conservative
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Matches 5; Conserv
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                                                                                                                                                                                                                                                                     A;Accession: G97084
A;Status: preliminary
A;Molecule type: DNA
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A,Gene: CAC1498
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C;Species: Staphylococcus aureus
C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004
C;Accession: H89992
R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; Oguc
C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.;
Lancet 357, 1225-1240, 2001
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A;Coss.references: UNIPROT:Q99SWB; GB:BA000018; PID:g13701812; PIDN:BAB43105.1; GSPDB:G
A;Experimental source: strain N315
C;Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        bombyxin A-1 homolog - ailanthus silkmoth
C;Species: Samia cynthia (ailanthus silkmoth)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
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                                                                                                                    25.0%; Score 5; DB 2; Length 98; 100.0%; Pred. No. 1.1e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A)Reference number: J00902
A)Accession: J00902
A)Molecule type: DNA
A)Residues: 1-99 </ri>
A)Residues: 1-99 </ri>
A)Cross-references: UNIPROT: P33718
A)Cross-references: Gesignated as Samia cynthia ricini
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R;Iwami, M.
submitted to JIPID, March 1991
                                                                                                              Query Match
Best Local Similarity 100.
Matches 5; Conservative
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Matches 5; Conserv
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Matches 5; Conserv
A;Gene: sbxA3
C;Superfamily: insulin
C;Keywords: hormone
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C,Superfamily: insulin
C,Keywords: hormone
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Genetics:

RESULT 45

ò g A; Gene: SA1825 Query Match

46

RESULT

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Gaps

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Length 117;

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Rivolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee, Joaly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium Clo A;Reference number: A96900; MUID:21359325; PMID:21359325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q97JT6; GB:AE001437; PIDN:AAK79159.1; PID:g15024109; GSPDB:GA;Experimental source: Clostridium acetobutylicum ATCC824
C;Genetics:
A;Gene: CAC1187
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(S.Species: Arabidopsis thaliana (mouse-ear cress)
(S.Species: Arabidopsis thaliana (mouse-ear cress)
(S.Species: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 27-Oct-2003
(S.Accession: T05357
(R.Bevan, M.) Terryn, N.; Ardiles, W.; Buysshaert, C.; Dasseville, R.; De Clerck, R.; De ewes, H.W.; Mayer, K.F.X.; Schueller, Submitted to the Protein Sequence Database, February 1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Species: Clostridium acetobutylicum
C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004
C;Accession: D97046
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A;Genome: nuclear
A;Gentrons: 32/1; 84/3; 100/3
A;Note: FBB4.170
C;Superfamily: ubiquinol-cytochrome-c reductase 14 kDa protein
C;Keywords: electron transfer; membrane protein; membrane-associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      [imported] - Clostridium acetobutylicum
                                    A;Gene: CPj0356
C;Superfamily: Chlamydia pneumoniae hypothetical protein CPn0356
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A, Accession: T05357
A, Molecule type: DNA
A, Residues: 1-122 cBEV-
A, Cross-references: EMBL-AL034567
A, Experimental source: cultivar Columbia, BAC clone F8B4
                                                                                                                                         25.0%; Score 5; DB 2; Len 100.0%; Pred. No. 1.3e+02; iive 0; Mismatches 0;
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100.0%; Pred. No. 1.3
iive 0; Mismatches
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Best Local Similarity 100...
S; Conservative
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Matches 5; Conservative
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Best Local Similarity
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   C, Genetics:
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A;Molecule type: DNA
A;Residues: 1-116 <KRM>
A;Residues: 1-116 <KRM>
A;Cross-references: UNIPROT:09Y8S4; DDBJ:AP000064; NID:g5105945; PIDN:BAA81576.1; PID:d1
C;Genetics: A;Gene: APE2559
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A;Cross-references: UNIPROT:Q92817; GB:BA000008; NID:g8978728; PIDN:BAA98564.1; GSPDB:GN
A;Experimental source: strain J138
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                                                                                                                                                                                                                                          25.0%; Score 5; DB 2; Length 116; 100.0%; Pred. No. 1.2e+02; ive 0; Mismatches 0; Indels
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Best Local Similarity
Matches 5; Conserv
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A; Title: Complete genome sequence of a multiple drug resistant Salmonella enterica serove A; Reference number: AB0502; MUD:21534947; PMID:11677608
A; Recession: AI0999
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-133 < PAR>A; Residues: 1-133 < PAR>A; Residues: 1-134 < PAR>A; Residues: 1-134 < PAR>A; Residues: 1-135 < PAR>A; Cross-references: GB:AL513382; PIDN:CAD08125.1; PID:g16505104; GSPDB:GN00176
C; Genetics:
A; Gene: yrfA
C; Superfamily: Escherichia coli hypothetical 16.9% protein b3392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein APE1600 - Aeropyrum pernix (strain K1)
C;Species: Aeropyrum pernix
C;Species: Aeropyrum pernix
C;Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
C;Accession: C7239
R;Kawarabayasi, Y.; Hino, Y.; Horikawa, H.; Yamazaki, S.; Haikawa, Y.; Jin-no, K.; Takahi awa, H.; Takamiya, M.; Masuda, S.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; Ki NNA Res. 6, 83-101, 1999
A;Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyria;Reference number: A72450; MUID:99310339; PMID:10382966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R'NG, W.V.; Kennedy, S.P.; Mahairas, G.G.; Berquist, B.; Pan, M.; Shukla, H.D.; Lasky, S.; Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, D.W.; Maddocks, D.G.; Jablor Jung, K.H.; Alam, M.; Freitas, T.

Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000
A;Authors: Hou, S., Daniels, C.J.; Dennis, R.P.; Omer, A.D.; Ebhardt, H.; Lowe, T.M.; Lië
A;Title: Genome sequence of Halobacterium species NRC-1.
A;Reference number: A84160; MUID:20504483; PMID:11016950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Residues: 1-133 <STO>
A;Cross-references: UNIPROT:Q9HSB7; GB:AE004437; NID:g10579938; PIDN:AAG18890.1; GSPDB:GR
C;Genetics:
A;Gene: VNG0311H
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C;Genetics:
A;Gene: APE1600
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C.Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
C.Accession: F84190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2; Length 133; . 1.4e+02;
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Best Local Similarity
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary A;Molecule type: DNA
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N'Alternate names: rev-like protein
('Species: caprine arthritis-encephalitis virus, CAEV
C'Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 09-Jul-2004
C'Accession: A40479
Kalinski, H.; Yaniv, A.; Mashiah, P.; Miki, T.; Tronick, S.R.; Gazit, A.
Virology 183, 786-792, 1991
A,Title: rev-like transcripts of caprine arthritis encephalitis virus.
A,Title: rev-like transcripts of caprine arthritis encephalitis virus.
A,Accession: A40479
A,Molecule type: mRNA
A,Molecule type: mRNA
A,Residences: UNIPROT:P31628; GB:M63105; NID:9323291; PIDN:AAA42893.1; PID:9323292
C,Genetics:
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Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.;
                                                                                                                                                             NyAlternate names: rev-like profein
C;Species: caprine arthritis-encephalitis virus, CAEV
C;Accession: F45343
R;Saltarelli, M; Querat, G; Konings, D.A.M.; Vigne, R.; Clements, J.E.
Virology 179, 347-344, 1990
A;Tille: Nucleotide sequence and transcriptional analysis of molecular clones of CAEV
A;Reference number: A45345; MUID:91021037; PMID:2171210
                                                                                                                                      trans-regulatory splicing-like protein - caprine arthritis-encephalitis virus (strain
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A;Gene: rev
A;Introns: 38/1
C;Superfamily: caprine arthritis-encephalitis virus rev-like protein
C;Superfamily: caprine arthritis-encephalitis virus rev-like protein
C;Keywords: splicing protein; transcription regulation
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C;Superfamlly: caprine arthritis-encephalitis virus rev-like protein
C;Keywords: splicing protein; transcription regulation
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A;Molecule type: mRNA
A;Residues: 1-133 <SAL>
A;Cross-references: UNIPROT:P33460; GB:M33677
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Matches
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T.L.; Hughes, A.L.; Bontrop, R.E.; Watkin
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C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
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A;Cross-references: UNIPROT:Q95533; EMBL:U05585; NID:g454787; PIDN:AAA50188.1; PID:g4547
C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 07-Jun.1996 #sequence_revision 07-Jun-1996 #text_change 09-Jul-2004
C;Accession: I38875
R;Garber, T.L.; Butler, L.M.; Trachtenberg, E.A.; Erlich, H.A.; Rickards, O. Immunogenetics 42, 19-27, 1995
Immunogenetics 42, 19-27, 1995
A;Fitle: HIA-B alleles of the Cayapa of Ecuador: new B39 and B15 alleles.
A;Reference number: I38860; MUID:95317819; PMID:7797264
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C,Species: Pan troglodytes (chimpanzee)
C,Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 09-Jul-2004
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C;Species: Pan troglodytes (chimpanzee)
C;Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 09-Jul-2004
C; Superfamily: class I histocompatibility antigen; immunoglobulin homology
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R;McAdam, S.N.; Boyson, J.E.; Liu, X.; Garber, T.L.; Hughes, A.L.;
Proc. Natl. Acad. Sci. U.S.A. 91, 5893-5897, 1994
A;Title: A uniquely high level of recombination at the HLA-B locus.
A;Reference number: IS9308; MUID:94286544; PMID:8016085
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R;McAdam, S.N.; Boyson, J.E.; Liu, X.; Garber, T.L.; Hughes, A.L.;
Proc. Natl. Acad. Sci. U.S.A. 91, 583-5897, 1994
A;Title: A uniquely high level of recombination at the HLA-B locus. A;Reference number: 159308; MUID:94286544; PMID:8016085
A;Accession: 180174
A;Staus: preliminary; translated from GB/EMBL/DDBJ
                                                                                        Length 137;
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                                                                            25.0%; Score 5; DB 2; Lenilarity 100.0%; Pred. No. 1.4e+02; Conservative 0; Mismatches 0;
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A,Molecule type: mRNA
A,Residues: 1-137 <RES>
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Best Local Similarity
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c;Species: Pan troglodytes (chimpanzee)
c;Date: 24-May-1996 #text_change 09-Jul-2004
c;Date: 24-May-1996 #text_change 09-Jul-2004
c;Date: 24-May-1996 #text_change 09-Jul-2004
c;Accession: 180173
R;MCAdam, S.N.; Boyson, J.E.; Liu, X.; Garber, T.L.; Hughes, A.L.; Bontrop, R.E.; Watkin Proc. Natl. Acad. Sci. U.S.A. 91, 5893-5897, 1994
A;Title: A uniquely high level of recombination at the HLA-B locus.
A;Reference number: 159308; MUID:94286544; PMID:8016085
A;Accession: 180173
A;Accession: 180173
A;Accession: Levil mary; translated from GB/BMBL/DDBJ
A;Accession: Lyce: DNA
A;Mesidues: 1-137 <RES>
A;Residues: 1-137 <RES>
A;Cross-references: UNIPROT:Q95532; EMBL:U05584; NID:g454785; PIDN:AAA50187.1; PID:g4547
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R;McAdam, S.N.; Boyson, J.E.; Liu, X.; Garber, T.L.; Hughes, A.L.; Bontrop, R.E.; Watkin Proc. Natl. Acad. Sci. Us.A. 91, 5893-5897, 1994
Proc. Natl. Acad. Sci. Us.A. 91, 5893-5897, 1994
A;Title: A uniquely high level of recombination at the HLA-B locus.
A;Reference number: 159308; MUID: 94286544; PMID: 8016085
A;Accession: 180175
A;Accession: 180175
A;Residue: preliminary; translated from GB/EMBL/DDBJ
A;Mclecule type: mRNA
A;Mcscules: 1-137 cRES
A;Residues: 1-137 cRES
A;Cross-references: UNIPROT: Q95534; EMBL: U05586; NID: 9454789; PIDN: AAA50189.1; PID: 94547
C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                        class I histocompatibility antigen - chimpanzee (fragment)
clspecies: Pan troglodytes (chimpanzee)
clspecies: Pan troglodytes (chimpanzee)
clspecies: Pan troglodytes (chimpanzee)
clspecies: Pan troglodytes (chimpanzee)
clspecies: 24-May-1996 #sequence_revision 24-May-1996 #text_change 09-Jul-2004
clspecies: 180172
Elect. Natl. Acad. Sci. Us.A. 91, 5893-5897, 1994
A.Title: A uniquely high level of recombination at the HLA-B locus.
A.Reference number: 159308; MUID: 94286544; PMID: 8016085
A.Accession: 180172
A.Accession: 180172
A.Accession: BOA
A.Residus: preliminary; translated from GB/EMBL/DDBJ
A.Residus: pr
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C;Species: Pan troglodytes (chimpanzee)
C;Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 09-Jul-2004
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Cispecies: Listeria monocytogenes
Cinate: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004
Cinate: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004
Cinate: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004
Ridlaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloecker,
Jones, L.M.; Rarst, U.
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; Maitournam, A.; Mat
ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; Voss, H.; Wehland,
A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
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A)Residues: 1-138 cGLAA>
A)Residues: 1-138 cGLAA>
A)Cross-references: UNIPROT:Q8Y976; GB:NC_003210; PIDN:CAC98735.1; PID:g16410046; GSPDB:CA)Experimental source: strain EGD-e
C;Genetics:
A;Gene: Imo0657
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   C;Accession: 138876
R;Garber, T.L.; Butler, L.M.; Trachtenberg, E.A.; Erlich, H.A.; Rickards, O.; De Stefano, Immunogentics 42, 19-27, 1995
Immunogentics 42, 19-27, 1995
Immunogentics 42, 19-27, 1995
A;Title HLA-B alleles of the Cayapa of Ecuador: new B39 and B15 alleles.
A;Reference number: 138860; MUID:95317819; PMID:7797264
A;Recession: 138876
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: preliminary; Aranslated from GB/EMBL/DDBJ
A;Residues: 1-137 <RES>
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A;Residues: 1-140 <BAR>
A;Cross-references: UMIPROT:Q9UT37; EMBL:AL121741; PIDN:CAB57336.1; GSPDB:GN00066; SPDB:£
A;Experimental source: strain 972h-; cosmid c824
C;Genetics:
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Matches 5; Conservative
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Matches 5; Conserv
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A;Status: preliminary
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A; Introns: 1/3; 17/3
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AI1156
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                                                                                                                             A,Cross-references: UNIPROT:Q29664; EMBL:U15639; NID:9930332; PIDN:AAA74046.1; PID:g9303
C,Superfamily: class I histocompatibility antigen; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A,Molecule_type: DNA
A,Residues: 1-137 <RES.
A,Cross-references: UNIPROT:Q29660; EMBL:U14756; NID:g930328; PIDN:AAC50171.1; PID:g9303
C,Superfamily: class I histocompatibility antigen; immunoglobulin homology
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R;Garber, T.L.; Butler, L.M.; Trachtenberg, E.A.; Erlich, H.A.; Rickards, O.; De Stefand Immunogenetics 42, 19-27, 1995
A;Title: HLA-B alleles of the Cayapa of Ecuador: new B39 and B15 alleles.
A;Reference number: I38860; MUID:95317819; PMID:7797264
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C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
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C;Species: Homo sapiens (man)
C;Date: 07-Jun-1996 #sequence_revision 07-Jun-1996 #text_change 09-Jul-2004
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C;Species: Homo sapiens (man)
C;Date: 07-Jun-1996 #sequence_revision 07-Jun-1996 #text_change 09-Jul-2004
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C;Species: Homo sapiens (man)
C;Date: 07-Jun-1996 #sequence_revision 07-Jun-1996 #text_change 09-Jul-2004
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                                 A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-137 <RES>
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A,Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local Similarity 100...
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A; Accession: I38875
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A;Residues: 1-152 <NRX.
A;Residues: 1-152 <NRX.
A;Cross-references: UNIPROT:095575; EMBL:AB030825; PIDN:BAA83159.1
A;Cross-references: UNIPROT:095575; EMBL:AB030825; PIDN:BAA83159.1
A;Experimental source: strain PAO1
B;Stower, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Br adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim, J. Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PAO1, an opportunistic patho A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: C83567
A;Status: preliminary
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C;Species: Archaeoglobus fulgidus
C;Species: Tober C; Accession: F695U9
F; Fleischmann, FD.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.; Fleischmann, K.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.; Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L.
Nature 390, 364-370, 1997
Forton, M.D.; Spriggs, T.; Artiach, F.; Kaine, B.P.; Sykes, S.
A,Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, F.; Kaine, B.P.; Sykes, S.
A,Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, F.; Kaine, B.P.; Sykes, S.
A,Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archaect
A,Accession: F695U9
A,Status: preliminary; nucleic acid sequence not shown; translation not shown
A,Mockeule type: DNA
A,Residues: 1-152 <a href="https://documbed.com/">MLES</a>
A,Cross-references: UNIPROT:028200; GB:AE000960; GB:AE00782; NID:g2689283; PIDN:AAB8918
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S, Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Barghar, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim.; Lory, S.; Olson, M.V.
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                                         T.; Kageyama, M.; Kanaya,
                              R;Nakayama, K.; Takashima, K.; Ishihara, H.; Shinomiya, T.; Kageyama, M.; Ka
submitted to the EMBL Data Library, August 1999
A;Description: Genetic relationship between bacteriocins and bacteriophages.
A;Reference number: Z22790
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100.0%; Pred. No. 1.5e+02;
.ive 0; Mismatches 0;
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C, Genetics:
A, Gene: PAO621
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C; Accession: T44544; C83567
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Best Local Similarity
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A; Residues: 1-152 <STO>
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c;Species: Lycopersicon esculentum (tomato)
c;Species: Lycopersicon esculentum (tomato)
c;Species: Lycopersicon esculentum (tomato)
c;Date: 22.Nov-1933 #sequence_revision 14-Jul-1995 #text_change 09-Jul-2004
c;Accession: 823801
R;Linthorst, H.J.M.; Danhash, N.; Brederode, F.T.; van Kan, J.A.L.; de Wit, P.J.G.M.; Bc A.L. inthorst, H.J.M.; Danhash, N.; Brederode, F.T.; van Kan, J.A.L.; de Wit, P.J.G.M.; Bc A.Fitle: Tobacco and tomato PR proteins homologus to win and pro-hevein lack the "hevein A;Reference number: 823799; MUID:92208317; PMID:1804403
A;Accession: 823801
A;Accession: 823801
A;Accession: 823801
A;Residues: L-143 <LINA
A;Residues: L-
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A.Note: this species has also been called Salmonella typhi
A.Note: this species has also been called Salmonella typhi
A.Note: this species has also been called Salmonella typhi
C.Accession: AE0812
B.P.Parkhill, J.; Dougan, G.; James, K.D.; Thomson, N.R.; Pickard, D.; Wain, J.; Churcher, F. T. Connerton, P.; Crohin, A.; Davis, R.M.; Pickard, D.; Wain, J.; Churcher, S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A.Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; A.Atchession: Reson as equence of a multiple drug resistant Salmonella enterica serov A, Reference number: AB0502; MUID:21534947; PMID:1167608
A.Status: preliminary
A,Nolecule type: DNA
A,Residus: 1-149 cPARA
A,Residus: 1-149 cPARA
A, Accession: Connected to the All Connected to
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A,Gene: STY2685
C,Superfamily: Bscherichia coli hypothetical protein b2433
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C;Date: 31-Jan-2000 #sequence_revision 31-Jan-2000 #text_change 09-Jul-2004
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25.0%; Score 5; DB 2; Length 140; 100.0%; Pred. No. 1.4e+02; cive 0; Mismatches 0; Indels
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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-157 <TIGR>
A;Cross-references: UNIPROT:P44005; GB:U32731; GB:L42023; NID:g3212193; PIDN:AAC22148.1;
C;Superfamily: hypothetical protein HI0489
                                                                             A.Accession: B83274
A.Status: preliminary
A.Status: preliminary
A.Molecule type: DNA
A.Status: Dradiminary
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A.Status: L154 < 2TO>
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A,Reference number: A82950; MUID:20437337; PMID:10984043
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C;Species: Haemophilus influenzae
C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 09-Jul-2004
C;Accession: E64008
R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Kerlavage, P.; Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kirkness, G.F.; Weidman, D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M. Science 269, 496-512, 1995
A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Reference number: A64000; MUID:95350630; PMID:7542800
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A,Baccession: AG0754
A,Accession: Delminary
A,Accession: Delminary
A,Residues: 1.156 < PAR>
A,Residues: 1.156 < PAR>
A,Cross-references: GB:AL513382; PIDN:CAD05738.1; PID:g16503230; GSPDB:GN00176
C,Genetics:
A,Gene: STY2199
C,Superfamily: very short patch repair endonuclease
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              GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                    1825181 segs, 575374646 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                       summaries
                                                                               - protein search, using sw model
                                                                                                                October 26, 2004, 06:48:06
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BAD17451
BAD17502
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BAC83842
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                                                                                                                                                                                                                                                                  60.0 , Gapext 60.0
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091NV7
091NV7
091NY1
091NY1
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Q913A2
Q88G85
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Q91319
Q913J0
Q77XS6
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Q6JQW3
Q6JQW4
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Q9YX82
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Q9INV9
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: uniprot_sprot:*
: uniprot_trembl:*
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length: 2000000000
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Maximum DB seq
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                                                                                                                                                                               Title:
Perfect so
Sequence:
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No.
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Q910H7

RESULT 1
0910H7
10 0910H7
10 0910
DT 01-M
DT 01-J-M
DT 0

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Mbayed V.A., Lopez J.L., Telenta P.F.S., Palacios G., Badia I., Ferro A., Galoppo C., Campos R.; Pirus genotypes in two different pediatric populations from Argentina ", J. Clin. Microbiol. 36:3362-3365(1998).

ENBL, AF043569, AAC79970.1; -.. GO. GO. 60:0016032; Priviral life cycle; IEA.

InterPro; PF000395; vMSA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

MEDLINE=98449868; PubMed=9774595;

MEDLINE=98449868; PubMed=9774595;

Mayed V.A., Lopez J.L., Telenta P.F.S., Palacios G., Badia I.,

Ferro A., Galoppo C., Campos R.;

"Distribution of hepatitis B virus genotypes in two different
pediatric populations from Argentina.";
J. Clin. Microbiol. 36:3362-3365 [1998).

EMBL, AF043565; AAC79966.1;

EMBL, AF043565; P.Viral life cycle; IEA.

InterPro; IPR00349; Hepvir_surfAg.

Pfam; PF00695; VMSA; 1.
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NEL TaxID=10407;
                                                                                                                                                                             Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus.
VCB__TaxID=10407;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      73 AA; 7961 MW; 194DE0279C879B1E CRC64;
                                                                             01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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MEDLINE=98449868; PubMed=9774595;
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                                                                                                                                            Surface antigen (Fragment).
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es 6; Conservative
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nes 6; Conservative
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                                     PRELIMINARY;
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                                                                                                                                                              Hepatitis B virus.
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NON TER
                   Q9WLQ2
ID Q9WLQ2
AC Q9WLQ2;
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Q9YXC7
RESULT 3
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STRAIR=ATCC 15692 / PAO1;
STRAINS=20437337; Pubmed=10984043;
Stover C.K., Pham X.-Q.T., Erwin A.L., Mizoguchi S.D., Warrener P.,
Hickey M.J., Brinkman F.S.L., Hufnagle W.O., Kowalik D.J., Lagrou M.,
Garber R.L., Goltry L., Tolentino E., Westbrock-Wadman S., Yuan Y.,
Brody L.L., Coulter S.N., Folger K.R., Kas A., Larbig K., Lim R.M.,
Smith K.A., Spencer D.H., Wong G.K.-S., Wu Z., Paulsen I.T.,
Reizer J., Saier M.H., Hancock R.E.W., Lory S., Olson M.V.;
"Complete genome sequence of Pseudomonas aeruginosa PAO1, an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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STRAIN=IG 63134.20;
Stuyver L., Van Geyt C., De Gendt S., Van Reybroeck G., Rossau R.;
Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                        Bacteria, Proteobacteria, Gammaproteobacteria, Pseudomonadales,
Pseudomonadaceae, Pseudomonas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HBsAg (Fragment).
Hepatitis B virus.
Viruses: Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 297;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR; C83311; C83311.
Complete proteome; Hypothetical protein.
SEQUENCE 297 AA; 33197 MW; 0DF94B92126D72D5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 1
56 AA; 5777 MW; 7C33FDDCD4B1F179 CRC64;
                                                                               01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-UUN-2003 (TrEMBLrel. 24, Last annotation update)
Hypothetical protein.
OrderedLocusNames=PA2661;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-00T-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
40.0%; Score 8; DB 2;
Best Local Similarity 100.0%; Pred. No. 0.57;
Matches 8; Conservative 0; Mismatches
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100.0%; Pred. No. 25;
tive 0; Mismatches
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                                         297 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           opportunistic pathogen.";
Nature 406:959-964(2000).
EMBL; AE004695; AAG06049.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                         PRELIMINARY;
                                                                                                                                                                                             Pseudomonas aeruginosa.
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Best Local Similarity
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                                                               Q910H7;
01-MAR-2001
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6VNI6Q

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Antigen.

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MEDLINE FROM N.A.

MEDLINE=2243060; PubMed=12534463;

MEDLINE=22423060; PubMed=12534463;

Melson K.E., Weinel C., Paulsen I.T., Dodson R.J., Hilbert H.,

Martins dos Santos V.A.P., Fouts D.E., Gill S.R., Pop M., Holmes M.,

Brinkac L.M., Beanan M.J., DeBoy R.T., Daugherty S.C., Kolonay J.F.,

Madupu R., Nelson W.C., White O., Peterson J.D., Khouri H.M.,

Hance I., Chris Lee P., Holtzapple E.K., Scanlan D., Tran K.,

Moazzez A., Utterback T.R., Rizzo M., Lee K., Kosack D., Moestl D.,

Wedler H., Lauber J., Stjepandic D., Hoheisel J., Straetz M., Heim S.,

Kiewitz C., Eisen J.A., Timmis K.N., Duesterhoeft A., Tuemmler B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pseudomonas putida (strain KT2440).
Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
              01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DDS-2001 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Complete genome sequence and comparative analysis of the metabolically versatile Pseudomonas putida KT2440."; Embrizon. Microbiol. 4:799-808(2002).
EMBL; AE016788; AAN69434.1; --
TIGR; PP3840; --
                                                                                                                                                                   Hijar Guerra G., Padilla Rojas C.P., Suarez Jara M.A., Cabezas Sanchez C.A.;
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
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73 73 73
73 AA; 7823 MW; 26A723DE136F0F8C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Complete proteome, Hypothetical protein.
SEQUENCE 74 AA, 8550 MW, 5F7FEBFB63468BD4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TrEMBLrel. 24, Created)
(TrEMBLrel. 24, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
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100.0%; Pred. No. 32;
iive 0; Mismatches
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100.0%; Pred. No. 32;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            74 A.A.
                                                                                                                                                                                                                      EMBL; AY059396; AAL26696.2; -...
GO; GO:0016032; P:viral life cycle; IEA.
InterPro; IPR000349; Hepvir_surfAg.
Pfam; P?00695; VMSA; 1.
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NCBI_TaxID=160488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical protein,
OrderedLocusNames=PP3840;
                                                                                                                                                                                                                                                                                                                                                                                                                     6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  47 SSWALG 52
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01-JUN-2003
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Surface antigen (Fragment).
Hepatitis B virus.
Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus.
                                                                                                                                                                  Hepatitis B Virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

Hijar Guerra G., Padilla Rojas C.P., Suarez Jara M.A.,
Cabezas Sanchez C.A.;
Submitted (ROV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AV053398; AAL26698.2;
GO, GO:0016032; P:viral life cycle; IEA.
InterPro; IFR000349; Heyvir_surfAg.
PF00695; vMSA; 1.
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73 73 73
73 AA, 7823 MW, 26A723DE136F0F8C CRC64;
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73 73
73 AA; 7923 MW; 26A723DE136F0F8C CRC64;
                                                                                           01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TrEMBLrel. 19, Created)
(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
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100.0%; Pred. No. 32;
tive 0; Mismatches
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100.0%; Pred. No. 32;
iive 0; Mismatches
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Pfam; PF00695; vMSA; 1.
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                                                                PRELIMINARY;
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                      NCBI_TaxID=10407;
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01-DEC-2001
01-DEC-2001
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Q913A2
ID Q9131
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Q913A1
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0; Indels

Length 73;

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Gaps

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0; Indels Length 74;

73 AA.

PRT;

PRELIMINARY;

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Surface antigen (Fragment).
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Matches 6; Conser
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                  Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10407;
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NON TER
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COYIFF
AC COYIF
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DT 01-MA
DT 01-JU
DE SULF
DE SULF
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Q77XS6
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MEDLINE=21642377; PubMed=11782923;

MEDLINE=21642377; PubMed=11782923;

Wan Steenbergen J.B., Niesters H.G.M., Op de Coul E.L.M.,

van Doornum G.J.J., Osterhaus A.D.M.E., Leentvaar-Kuijpers A.,

Coutinho R.A., van den Hoek J.A.R.;

"Molecular epidemiology of hepatitis B virus in Amsterdam 1992-1997.";

J. Med. Virol. 66:159-165(2002).

GO: GO:0016032; Priral life cycle; IEA.
                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=ESV-1;
STRAIN=ESV-1;
Bolaroque N., Bothe G., Pohl T., Knippers R., Mueller D.G., Boland W.;
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF204951; AAK14633.1; -.
SEQUENCE 77 AA; 9075 MW; 2D7B8BF96D113B2C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                Ectocarpus siliculosus virus.
Viruses; dsDNA viruses, no RNA stage; Phycodnaviridae; Phaeovirus.
11.
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NCBI_TaxID=10407;
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9070 MW; 0532646A3D2D4847 CRC64;
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Last annotation update)
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(TremBirel. 19, Last sequence update)
(TremBirel. 24, Last annotation update)
                                                                                          Last sequence update)
Last annotation update)
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100.0%; Pred. No. 30,
0; Mismatches
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Pfam; PF00695; vMSA; 1.
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                                                                         01-JUN-2002 (TrEMBLrel. 21,
01-JUN-2002 (TrEMBLrel. 21,
01-JUN-2002 (TrEMBLrel. 21,
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Best Local Similarity 100.
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Matches 6; Conservative
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82
82 AA;
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01-JUN-2003 (
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Q913J0;
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Q913I9;
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Q913J0
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Q8QN71
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                                                                                         MEDLINE=21642377; PubMed=11782923;
MEDLINE=21642377; PubMed=11782923;
MEDLINE=21642377; PubMed=11782923;
Van Steenbergen J.E., Niesters H.G.M., Op de Coul E.L.M.,
van Doornum G.J.,J., Osterbaus A.D.M.E., Leentvaar-Kuijpers A.,
Coutinho R.A., van den Hoek J.A.R.;
"Moleoular epidemiology of hepatitis B virus in Amsterdam 1992-1997.";
J. Med. Virol. 66:159-165(2002).

EMBL; AY048640; AAL06221.1;
GO, GO:016032; P:Viral life cycle; IEA.

InterPro; IPR00349; Hepvir_surfAg.
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STRAIN=2997AIN=2997.
STRAIN=2974.
Van Geyt C., De Gendt S., Rombout A., Wyseur A., Maertens G.,
Rossau R., Stuyver L.;
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
BEMBL, AF045010; AAD02372.1;
NON_TER
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
[1]
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       %.70L-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 83 AA; 9183 MW; 082532646A3D2D48 CRC64;
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100.0%; Pred. No. 35;
iive 0; Mismatches
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100.0%; Pred. No. 41;
tive 0; Mismatches
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STRAIN=22298;
Van Geyt C., De Gendt S., Rombout A., Wyseur A., Maertens G., Rossau R., Stuyver L.;
Submitted (JAN.1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF045008; AAD02370.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=29165;
Van Geyt C., De Gendt S., Rombout A., Wyseur A., Maertens G., Rossau R., Stuyver L.;
Rossau R., (JAN-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis Virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-UNY-2093 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
                                                                                                                                                                                                                    0; Indels
     Rossau R., Stuyver L.;
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF045012; AAD02374.1; -
GO; GO:0016032; P:viral life cycle; IEA.
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
Antigen.
NON_TER 1 1 1
NON_TER 98 88
SEQÜENCE 98 AA; 10887 MW; COBBCD7D9187ASAE CRC64;
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                                                                                                                                                                                    Length 98;
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
                                                                                                                                                                                 30.0%; Score 6; DB 2;
100.0%; Pred. No. 41;
ative 0; Mismatches
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100.0%; Pred. No. 41;
tive 0; Mismatches
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                STRAIN=29976;
Van Geyt C., De Gendt S., Rombout A., Wyseur A., Maertens G., Rossau R., Stuyver L.;
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF045011; AAD03373.1;
GO; GO:0016032; P:viral life cycle; IEA.
InterPro; IPR000349; Hepvir_surfAg.
PF00695; VMSA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=29981;
Van Geyt C., De Gendt S., Rombout A., Wyseur A., Maertens G., Rossau R., Stuyver L., De Gendt S., Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF045013; AAD02375.1; --
GO; GO:0015032; P:Viral life cycle; IEA.
InterPro; IPR00049; Hepvir_surfAg.
PF00695; VMSA; 1.
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Van Geyt C., De Gendt S., Rombout A., Wyseur A., Maertens G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-UNY-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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100.0%; Pred. No. 41;
tive 0; Mismatches 0; Indels
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98 98
98 AA; 10901 MW; 1CFBD068D337A646 CRC64;
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98 98
98 AA; 10899 MW; E7704BEA5192E1BB CRC64;
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01-WAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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Surface antigen (Fragment).
Hepatitis B virus.
                                                                                                                                                                                                                                           Best_Local Similarity 100.
Matches 6; Conservative
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STRAIN=IG 63135.20;
Survare L., Van Geyt C., De Gendt S., Van Reybroeck G., Rossau R.;
Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF102608; AAF80717.1; -.
EMBL; AF102608; AAF80717.1; -.
InterPro; IPR000349; Hepvir_surfAg.
                                                                                                                                                                                                                                                                                                                                                        Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
VCBI_TaxID=10407;
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99 AA, 11112 MW, DD4F202E6FB1E5E1 CRC64;
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99 99
99 AA, 11040 MW; DD4F202CCD3BCFEI CRC64;
99 99 AA; 11112 MW; DD4F202E6FB1E5E1 CRC64;
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01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UTD-2003 (TrEMBLrel. 24, Last annotation update)
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01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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                                               30.0%; Score 6; DB 2;
100.0%; Pred. No. 41;
tive 0; Mismatches
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100.0%; Pred. No. 41;
cive 0; Mismatches
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Q9INV7
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STRAIN=[66 3138.20;
Stuyver L., Van Gert C., De Gendt S., Van Reybroeck G., Rossau R.;
submitted (ocr-1998) to the EMBL/GenBank/DDBJ databases.
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
HBSAG (Fragment)
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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Stuyver L., Van Gerdt S., Van Reybroeck G.,
Stuyver L., Van Gert C., De Gendt S., Van Reybroeck G.,
Submitted (CCT-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, AF102597; AAF80695.1; -- GO, GO, GO, GO, P. VITAB 1 life cycle; IEA.
InterPro; IPR00349; Heyvir_surfAg.
Pfam; PF00695; VMSA; 1.
                                                                                                                                           Length 98;
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98 98
98 AA; 10758 MW; EA69315B0AECF8D4 CRC64;
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98 98
98 AA; 10878 MW; 01B55868C192FABE CRC64;
                                                                                                                                                                                                                                                                                                                                                                        01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                             DB 2;
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   GO:0016032; P:viral life cycle; IEA.
   GO, GO:0016032, P:viral life cycle;
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
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Matches 6; Conserv
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Best Local Similarity
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Q9INV5
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001NX3
001NX
AC 091NX
DT 01-0C
DT 01-0C
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DE HEBAG
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Gaps

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Length 99;

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STRAIN=NCC 2705;
MREDLINE=22294977; PubMed=12381787;
SCHOLI M.A., Karmirantzou M., Snel B., Vilanova D., Berger B., Schell M.A., Karmirantzou M., Snel B., Dessi G., Zwahlen M.-C., Desiere F., Bork P., Delley M., Pridmore R.D., Arigoni F.;
The genome sequence of Bifidobacterium longum reflects its adaptation to the human gastrointestinal tract.";
Proc. Natl. Acad. Sci. U.S. A. 99:14422-14427(2002).
GO; GO:0016020; C:membrane; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oryza sativa (japonica cultivar-group).
Bukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae, Oryzeae; Oryza, Oryza sativa.
                                                                                                                                                                                                                                                                                                        Bacteria, Actinobacteria, Actinobacteridae, Bifidobacteriales,
Bifidobacteriaceae, Bifidobacterium.
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STRANBEAU. Nipponbare;
Sasaki T., Matsunoto T., Katayose Y.;
"Oryza sativa nipponbare(GA3) genomic DNA, chromosome 2, BAC
clone:OSUMBD046012.",
Submitted (AUG-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AP005643; BAD17451.1; -.
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100.0%; Pred. No. 41;
live 0; Mismatches 0; Indels
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Pfam, PR02325; YGGT; 1.
Complete proteome; Hypothetical protein.
SEQUENCE 100 AA; 11654 MW; 7B219E7C9B016BF9 CRC64;
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100 Aa; 10960 MW; CC21D0B5AFBD0E4D CRC64;
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Hypothetical protein OSJNBb0046012.18.
Orvan
                                                                                                                                                                   (TrEMBLrel. 23, Last sequence update) (TrEMBLrel. 24, Last annotation update)
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                                                                                      100 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30.0%; Score 6; DB 2
Similarity 100.0%; Pred. No. 41;
6; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-WAY-2004 (TrEMBLrel. 27, Created)
10-WAY-2004 (TrEMBLrel. 27, Last seq
10-WAY-2004 (TrEMBLrel. 27, Last ann
                                                                                                                                            Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                      PRT;
                                                                                                                                          23,
                                                                                                                                                                                                                                                       OrderedLocusNames=BL0120;
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                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                            (TrEMBLrel.
                                                                                                                                                                                                                                                                              Bifidobacterium longum
                                                                                                                                                                                                                       Hypothetical protein
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Best Local Similarity
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Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11 RWLRRY 16
                                                                                                                                                                                                                                                                                                                                                                 NCBI TaxID=216816;
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   62 RWLRRY 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 WRWLRR 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WRWLRR 24
                                                                                                                                       01-MAR-2003
01-MAR-2003
01-JUN-2003
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BAD17451;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                Q8G7W5;
                                                                                      08G7W5
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                               RESULT 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 25
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                                                            08G7W5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN=ICCG 3813;
Strain=ICCG 3813;
Stuyver L., Van Gerdt S., Van Reybroeck G., Rossau R.;
Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF102596; AAF80693.1;
EMBL; Printal life cycle; IEA.
InterPro; IPR00349; Hepvir_surfAg.
PF00695; vMSA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oryza sativa (japonica cultivar-group).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus.
NCBI_TaxID=10407;
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  Indels
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Submitted (AUG-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AP005644; BAD17502.1;
EMBL, AP005644; BAD17451.1;
Hypothetical protein.
SEQUENCE 100 AA; 10960 MW; CC21D0B5AFBD0E4D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 99;
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99 99
99 AA; 11004 MW; BF67704BEF4592E1 CRC64;
                                                                                                                                                                                                                                                                                               01-0cT-2000 (TrEMBLrel. 15, Created)
01-0cT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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                                                                                                                                                                                                                                                 99 AA.
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100.0%; Pred. No. 41;
tive 0; Mismatches
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  Mismatches
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6; Conservative
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                       46 SSWALG 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 WRWLRR 15
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                                                   4 SSWALG 9
                                                                                                                                                                                                                                                                                                                                                                                  HBsAg (Fragment).
Hepatitis B virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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NON TER
SEQUENCE
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Matches
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Gaps

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Length 100;

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PubMed=15184419;
Stink R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Da Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian
                        Oryza sativa (japonica cultivar-group).
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Ehrhartoideae; Oryzeae; Oryza; Oryza sativa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S protein (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae, Orthohepadnavirus.
                                                                                                                                                 STRAIN=cv. Nipponbare;
Sasaki T., Matsumoto T., Katayose Y.;
"Oryza sativa nipponbare(GA3) genomic DNA, chromosome 7, BAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S protein (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
                                                                                                                                                                                                                                                                                                   Length 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 113;
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                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                             Close Columbotoning to the EMBL/GenBank/DDBJ databases. Submitted (MAX-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AP005172; BAC83842.1; -. Hypothetical protein. SEQUENCE 101 AA; 10261 NW; 238F8F70BF4D8092 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  113 AA; 12242 MW; FIB6DA4F0D5E50E2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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                                                                                                                                                                                                                                                                                                   DB 2;
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                                                                                                                                                                                                                                                                                                                     100.0%; Pred. No. 42; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          113 AA
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100.0%; Pred. No. 46;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              patients.";
J. Clin. Microbiol. 42:2455-2460(2004).
EMBL; AA233405; AAQ97022.1;
INTERFO; IPRO00349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
Hypothetical protein OSJNBb0002J01.11.
                                                                                                                                                                                                                                                                                                     Score 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                     30.0%;
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Matches 6; Conservative
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                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10407;
                 OSJNBB0002J01.11.
                                                                                                                                                                                                                                                                                                                                                                                                       66 WRWLRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q6JQW1;
05-JUL-2004
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NON TER
SEQUENCE
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Q6JQW3
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                                                                                                                                                               Oryza sativa (japonica cultivar-group).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza; Oryza sativa.
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Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
Ehrhartoideae, Oryzeae, Oryza.
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                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=cv. Nipponbare;
Sasaki T., Matsumoto T., Katayose Y.;
Sasaki T., Matsumoto T., Katayose Y.;
Susaki T., Matsumoto T., Matsumoto T., Matsumoto T., BAC
clone:OSJNBD0056122.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2; Length 101;
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Sasaki T., Matsumoto T., Katayose Y.;
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AP005172; BAC83842.1; -
Hypothetical protein.
SEQUENCE 101 AA; 10261 MW; 238F8F70BF4D8092 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 100
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                                                                                                                                                                                                                                                                                                                                                                                         al protein.
100 AA; 10960 MW; CC21D0B5AFBD0E4D CRC64;
                                                                                LU-MAY-2004 (TERMELTEL. 27, Last sequence update)
Hypothetical protein OSJNBb0056122.58.
OSJNBB0056122.58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       30.0%; Score 6; DB 2
100.0%; Pred. No. 41;
tive 0; Mismatches
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100.0%; Pred. No. 42;
tive 0; Mismatches
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Hypothetical protein OSJNBb0002J01.11
Name-OSJNBb0002J01.11;
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                                                    PRT;
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es 6; Conservative
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Best Local Similarity
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02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                         Hypothetical
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BAC83842;
                                                  BAD17502
BAD17502;
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BAC83842
                   RESULT 26
BAD17502
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PubMed=15184419;
Sitnik R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Da Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian
patients.";
J. Clin. Microbiol. 42:2455-2460(2004).
EMBL; AY323401; AAQ97018.1;
InterPro; IPR000349; Hepvir_surfAg.
Pfou695; vMSA; 1.
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PubMed=15.84419;
Sitnik R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Da Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian patients.";
J. Clin. Microbiol. 42:2455-2460(2004).
EMBL, FA323399, AAQ97016.1;
InterPro; IPR000349; Hepvir_surfAg.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
5-JUL-2004 (TrEMBLrel. 27, Last annotation update)
5 protein (Fragment).
7 protein B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBL_TaxID=10407;
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113 113
113 AA; 12273 MW; F1A39B1B1AF32A22 CRC64;
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113 113
113 AA; 12256 MW; 86B6DA4F0D5E55E2 CRC64;
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05-UUL-2004 (TrEMBLrel. 27, Created)
05-UUL-2004 (TrEMBLrel. 27, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
S protein (Fragment).
Hepatitis B virus.
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100.0%; Pred. No. 46;
tive 0; Mismatches
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les 6; Conserv
[1]
SEQUENCE FROM N.A.
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Best Local Similarity
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SEQUENCE
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NON TER
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QGJQW8;
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Matches
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                                                                                                      Sitnik R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Da Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian
patients.";
J. Clin. Microbiol. 42:2455-2460(2004).
EMBL, PA323403, AAQ97020.1;
InterPro; IPR000349; Hepvir_surfAg.
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Da Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian
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Hepatitis B virus.

Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.

NCBI_TaxID=10407;
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NCBI_TaxID=10407;
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113 113
113 AA; 12242 MW, F1B6DA4F0D5E50E2 CRC64;
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113 113
113 AA; 12242 MW; FIB6DA4F0D5E50E2 CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEWBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
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100.0%; Pred. No. 46;
tive 0; Mismatches
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J. Clin. Microbiol. 42:2455-2460(2004).
EMBL; AX323402; AAQ97019.1; -.
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Pfam; PF00695; vMSA; 1.
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Best Local Similarity
Matches 6; Conserv
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SEQUENCE FROM N.A.
PubMed=15184419;
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Best Local Similarity
Matches 6; Conserv
                                                   SEQUENCE FROM N.A.
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NCBI_TaxID=10407;
                                                                              PubMed=15184419;
                                                                                                                                                                                                                                                                                                                                               Antigen.
NON TER
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Q6JQW4
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"Hepatitis B virus genotypes and precore and core mutants in brazilian
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NCBL_raxID=10407;
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NCBI_TaxID=10407;
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STRAIN=072 96;
Lyra A.C., Pinho J.R.R., Mello I.M.V.G.C., Carrilho F.J.,
da 311va J.C.;
Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AY133290; AAN15315.1; -.
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Lyra A.C., Pinho J.R.R., Mello I.M.V.G.C., Carrilho F.J., da Silva L.C.;
Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AV133291, ANN15316.1, -
GO, GO:0016032; Prviral life cycle; IEA.
InterPro; IRR000349; Hepvir_surfAg.
                                                                                                                                                                                                                            Length 113;
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113 AA, 12242 MW, FIB6DA4F0D5E50E2 CRC64;
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119 AA; 13015 MW; A44C2CD2C28B5C77 CRC64;
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-UJN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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100.0%; Pred. No. 46;
rative 0; Mismatches
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100.0%; Pred. No. 48;
tive 0; Mismatches
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                       patients.";
J. Clin. Microbiol. 42:2455-2460(2004).
BMBL, 743233396, AAQ97013.1;
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
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Hepatitis B virus.
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Matches 6; Conservative
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PubMed=15184419;
Sitnik R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Ba Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian patients.";
J. Clin. Microbiol. 42.2455-2460(2004).

EMBL; AY323397; AAQ97014.1;
InterPro, IPR000349; Hepvir_surfAg.

Pfam; PF00695; vMSA; 1.
                       PubMed=15184419;
Sitnik R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Da Silva L.C., Carrilho F.J.;
"Hepatitis B virus genotypes and precore and core mutants in brazilian
patients.";
J. Clin. Microbiol. 42:2455-2460(2004).
BMBL; Microbiol. 42:2455-2460(2004).
NON TER 1.13 1.13
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Sitnik R., Rebello Pinho J.R., Bertolini D.A., Bernardini A.P.,
Da Silva L.C., Carrilho F.J.;
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NCBI_TaxID=10407;
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Hepatitis B virus.
Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus.
NCBI_TaxID=10407;
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113 AA; 12242 MW; F1B6DA4F0D5E50E2 CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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100.0%; Pred. No. 46;
iive 0; Mismatches
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100.0%; Pred. No. 46;
Live 0; Mismatches
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Best Local Similarity 100.0
Matches 6; Conservative
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Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O., Carballal G., Quarleri J.F., D'Wello F., Howard C.R., Oubina J.R.; "Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies."; J. Med. Virol. 69:18-26 (2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
MEDIATE=223367; PubMed=12436473;
MATHRE=223367; PubMed=12436473;
MATHRE V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepatitis B virus S gene mutants in a patient with chronic active
hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
                                                                            SEQUENCE FROM N.A.

Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
Oubina J.;
Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.

EMBL, AF500256; AAM27436.1; -.
GO, GO:0016032; P:viral life cycle; IEA.

InterPro; IPR000349; Hepvir_surfAg.
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
Oubina J.;
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EMBL, AFS00255; AM27435.1; -.
GO; GO:0014632; P:Viral life cycle; IEA.
InterPro; IPR000349; Hepvir_surfAg.
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119 119
119 AA, 13088 MW, 103DBAC4DAB0A750 CRC64;
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100.0%; Pred. No. 48;
tive 0; Mismatches
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ive 0; Mismatches
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Hepatitis B virus.
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Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
Oubina J.;
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EMBL; AF500257; AAM27437.1; -...

GO; GO:0016032; P:viral life cycle; IEA.

Interpro; IFR000349; Heyvir_surfAg.

Pfam; PF00695; VMSA; 1.
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 GO:0016032; P:viral life cycle; IEA
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01-OCT-2002 (TrEMBLrel. 22,
01-OCT-2002 (TrEMBLrel. 22,
01-JUN-2003 (TrEMBLrel. 24,
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01-OCT-2002 (TrEMBLrel. 22,
01-OCT-2002 (TrEMBLrel. 22,
01-JUN-2003_ (TrEMBLrel. 24,
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Les 6; Conservative
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Hepatitis B virus.
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Hepatitis B virus.
               InterPro; IPR000349;
Pfam; PF00695; vMSA;
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MEDILTE=22233367; PubMed=12436473;
MATHOE V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepartitis B virus S gene mutants in a patient with chronic active
hepartitis with dirculating mati-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
Oubina J.;
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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NCBI_TaxID=10407;
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Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.
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119 AA; 13148 MW; 174DBDA6A51A1351 CRC64;
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                     01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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100.0%; Pred. No. 48;
tive 0; Mismatches
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GO; GO:0016032; P: Viral life cycle; IEA.
InterPro; IPR000349; Hepvir_surfag.
Pfam; PF00695; vMSA; 1.
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MEDILINE-22233367, PubMed=12436473;
MATHEL V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepatitis B virus S gene mutants in a patient with chronic active
hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
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MEDLINE-223367, PubMed=12436473;

Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,

Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;

Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies.";

J. Med. Virol. 69:18-26(2003).
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
Oubina J.;
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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Submitted (APR-2002) to the BMBL/GenBank/DDBJ databases.
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119 119 119 AA, 13251 MW; 8A4D6C1BB5066D2A CRC64;
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MEDLINE=22333367; PubMed=12436473;
MAIDLE=22333367; PubMed=12436473;
MAIDLE M. . Bepinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
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NCBI_TaxID=10407;
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Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AFS00250; AMM27430.1; --
GO; GO:0016032; P:viral life cycle; IBA.
InterPro; IPR000349; Hepvir_surfAg.
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                                                                         119 AA; 13071 MW; 22387D0FB6736BF9 CRC64;
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GO; GO:0016032; P:V1ral life cycle; IEA.
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                                     Pfam; PF00695; vMSA; 1
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Hepatitis B virus.
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SEQUENCE FROM N.A. MBDMed=12436473; MBDLINE=22333367; PubMed=12436473; Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O., Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.; "Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating anti-HBs antibodies."; Med. Virol. 69:18-26(2003) A.
                                              Mando 0.,
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MEDLINE=22333367, PubMed=12436473,
Mathet V.L., Fald M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.,
"Hepatitis B virus S gene mutants in a patient with chronic active
hepatitis with circulating Anti-HBs antibodies.";
"Med. Virol. 69:18-26(2003).
                                                                                                                                                                SEQUENCE FROM N.A.
Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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NCBI_TaxID=10407;
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Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.
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Last annotation update)
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GO; GO:0016032; P:viral life cycle; IEA.
InterPro; IPR00349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
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MEDLINE=22323367; PubMed=12436473; Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O., Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.; Carballals G., Quarneri J.F., D'Mello F., Howard C.R., Oubina J.R.; Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies."; d. Med. Virol. 69:18-26(2003).
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SECUENCE FROM N.A.

Madblines_2333367; PubMed=12416473;

Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,

Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;

Carballal G. Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;

Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies.";

J. Med. Virol. 69:18-26(2003).
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus.
NCBI_TaxID=10407;
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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ive 0; Mismatches
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GO; GO:0016032; P:01ral life cycle; IEA.
InterProy. IPR000349; Hepvir_surfAg.
Pfam; PF00695; VMSA; 1.
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Best Local Similarity 100.
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                                                                                                                                                  SEQUENCE FROM N.A.

MEDLINE=22323367; PubMed=12436473;
Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepatitis B virus S gene mutants in a patient with chronic active
hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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NCBI_TaxID=10407;
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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EMBL, AFS00243; AAM27423.1; -.
GO, GO:0016032; P:viral life cycle; IEA.
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GO, GO:0016032; P:viral life cycle; IEA.
InterPro; IPR000349; Hepvir surfAg.
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Hepatitis B virus.
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MEDLINE-2233367; PubMed=12436473;
MEDLINE-2233367; PubMed=12436473;
MATHET V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O.,
Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Oubina J.R.;
"Hepatitis B virus S gene mutents in a patient with chronic active
hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
                                                     Mando 0.,
                                       MEDINE=22323367; PubMed=12436473; Sanchez D.O., Ruiz V., Mando O., Mathet V.L., Feld M., Espinola L., Sanchez D.O., Ruiz V., Mando O., Carballal G., Quarleri J.F., D'Mello F., Howard C.R., Cubina J.R., "Hepatitis B virus S gene mutants in a patient with chronic active hepatitis with circulating Anti-HBs antibodies.";
J. Med. Virol. 69:18-26(2003).
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
Oubina J.;
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
S antigen (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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EMBL; AFS00237; AAM27417.1; -.
GO; GO:0016032; P:viral life cycle; IEA.
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119 119
119 AA; 13114 MW; 1A2DB0CBC5177351 CRC64;
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119 AA; 13148 MW; 103DBAD1D51A1354 CRC64;
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100.0%; Pred. No. 48;
cive 0; Mismatches
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100.0%; Pred. No. 48;
ive 0; Mismatches
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NCBI_TaxID=10407;
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 Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J.,
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Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri
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NCBL_TaxID=10407;
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                           (APR-2002) to the EMBL/GenBank/DDBJ databases.
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Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases
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119 AA; 13285 MW; E53B101764C7C855 CRC64;
                                                                                                                                       119 AA; 13278 MW; 1758A8A6A50679E1 CRC64;
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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100.0%; Pred. No. 48;
tive 0; Mismatches
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                                        EMBL; AF500241; AAM27421.1; -.
GO; GO:0016032; P:viral life cycle; IEA
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                                                                  InterPro, IPR000349; Hepvir surfAg
Pfam; PF00695; vMSA; 1.
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nes 6; Conservative
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01-OCT-2002 (TrEMBLrel.
01-JUN-2003 (TrEMBLrel.
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Hepatitis B virus.
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QBJYY6

RESULT 56 **OBJYY6** 

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MEDLINE=91276235; PubMed=2055466;
Yamazoe M., Nakai S., Ogasawara N., Yoshikawa H.;
Yamazoe M., Nakai S., Ogasawara N., Yoshikawa H.;
Yamazoe M., Nakai S., Ogasawara N., Yoshikawa H.;
Sites (v-k and gag-1ike) in a hepaticelluar carcinoma.";
Gene 100:139-146(1991).
BMBL; M60766; AAA37108.1; -.
BMBL; M60766; AAA37108.1; -.
GO, GO:0016032; P:viral life cycle; IEA.
InterPro; IPR000349; Hepvir_surfAg.
Ffam; PR00692; WGA, 1.
NON TER 123 A2, 14168 MW; 56196675B7CZAD90 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-NC89-15;
Arankalle V.A., Murhekar K.M., Gandhe S.S., Murhekar M.V.,
Ramdasi A.Y., Padbidri V.S., Sehagal S.C.;
"Hepatitis B virus: Predominance of Genotype D in hyper-endemic
primitive tribes of Andaman and Nicobar Islands, India (1989-1999).";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY155433; AA017894.1; -.
NON TER 121
NON TER 121
SEQÜENCE 121 AA; 13138 WW; C18EBB3C94400B1B CRC64;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Sciuridae; Sciurinae;
                                                                                                                                                     AAO17894;
02-MAR-2004 (TEEMBLEE]. 27, Created)
02-MAR-2004 (TEEMBLEE]. 27, Last sequence update)
02-MAR-2004 (TEEMBLEE]. 27, Last annotation update)
Surface antigen (Fragment).
Wiruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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123 AA; 14168 MW; 56196675B7C2AD90 CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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100.0%; Pred. No. 48;
iive 0; Mismatches
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100.0%; Pred. No. 49;
iive 0; Mismatches
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111 SSWALG 116
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Best Local Similarity
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NCBI_TaxID=9995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. Mathet V., Feld M., Ruiz V., Espinola L., Sanchez D., Quarleri J., Oubina J.; Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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121 AA; 13138 MW; C18EBB3C94400B1B CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Surface antigen (Fragment).
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100.0%; Pred. No. 48;
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100.0%; Pred. No. 48;
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01-0CT-2002 (TrEMBLrel. 22,
01-JUN-2003 (TrEMBLrel. 24,
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Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best_Local Similarity 100.
Matches 6; Conservative
                                                                          PRELIMINARY;
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                                                                                                                                                                                                                                                                                       NCBI_TaxID=10407;
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Q6YF40

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RESULT 57 Q6YF40

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01-FEB-1995
01-FEB-1995
05-JUL-2004
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Takahashi M., Nishizawa T., Gotanda Y., Tsuda F., Komatsu F.,
Takahashi M., Nishizawa T., Altankhuu M., Chimedregzen U., Narantuya L.,
Rawabata T., Hasegawa K., Kagawa Y., Okamoto H.;
"High prevalence of antibodies to hepatitis A and E viruses and
viremia of hepatitis B, C, and D viruses among apparently healthy
populations in Mongolia";
Clin. Diagn. Lab. Immunol. 11:392-398(2004).
EMBL; AB119015; BAD12500.1;
                                                                                                                                                                                                                                                                                                                                 Takahashi M., Nishizawa T., Gotanda Y., Tsuda F., Komatsu F.,
Takahashi M., Hasegawa K., Altankhuu M., Chimedregzen U., Narantuya L.,
Hoshino H., Hino K., Kagawa Y., Okamoto H.,
High prevalence of antibodies to hepatitis A and E viruses and
viremia of hepatitis B, C, and D viruses among apparently healthy
populations in Mongolia."; 11:392-398(2004).
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus
                                                                                                                                                                                                             Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30.0%; Score 6; DB 2; Length 132; 100.0%; Pred. No. 52; 0; Indels iive 0; Mismatches 0; Indels
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132 132
132 AA, 14320 MW, A076867777BF7864 CRC64;
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24-MAR-2004 (TrEMBLrel. 27, Last sequence update)
24-MAR-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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                                               132 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; ABI19015; BAD12500.1; -.
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
                                                  PRT;
                                                                                                                                                                Surface antigen (Fragment).
Hepatitis B virus.
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                                               PRELIMINARY;
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Matches 6; Conserv
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                    NCBI_TaxID=10407;
                                                                                                                                                                                                                                                                                                                  PubMed=15013993;
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24-MAR-2004
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RESULT 60
Q75ZV2
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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TISSUE-4-day old flower tepal;
TISSUE-4-day old flower tepal;
THEATIFICATION of genes associated with perianth senescence in affectification of genes associated with perianth senescence in affection of genes pseudomarcissus L. 'Dutch Master').";
Then Sci. 163:13-21(2002).
The Plant Sci. 163:13-21(2002).
The Plant Sci. 163:13-21(2002).
The REBL; AF462230; AAL63931.; -..
THEATING CONDITION INTEGRAL TOWN THEATING CONDITION OF CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-2002 (TrEWBLrel. 20, Last sequence update)
01-JUN-2003 (TrEWBLrel. 24, Last annotation update)
Transporter-like protein (Fragment)
Narcissus pseudonarcissus (Daffodil).
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Amaryllidaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                        Name=US426;
Gallid herpesvirus 1 (strain GA) (GaHV-1) (Marek's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein.
SEQUENCE 142 AA; 15662 MW; 76D137DF02735E26 CRC64;
                                                                                                                                                                                                                                                                                                                 Virūses; dsDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Marek's disease-like viruses.
NCBI_TaxID=10388;
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                                                                                                            Last sequence update)
Last annotation update)
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100.0%; Pred. No. 55;
.ive 0; Mismatches
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142 AA.
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                                                                         (Rel. 31, Created)
                                                                                                                                           05-JUL-2004 (Rel. 44, Last ann
Hypothetical 15.7 kDa protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; M80595; AAB59898.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30.08;
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Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.
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Matches 6; Conserval
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                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                15 SSWALG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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SEQUENCE FROM N.A.
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NON TER
SEQUENCE
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Q6W5E6
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AAQ86848
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tanigami A., Fujiwara T., Shibahara T., Goto Y., Hirao M., Shimizu F., Wakebe H., Ono T., Hishigaki H., Watanabe T., Ozaki K., Sugiyama T., Irie R., Otsuki T., Sato H., Wakamatsu A., Ishihi S., Yamamoto J., Isono Y., Kawai-Hio Y., Satto K., Nishikawa T., Kimura K., Yamashita H., Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K., Wagatsuma M., Murakawa K., Kanehori K., Takahashi-Fujii A., Oshima A., Sugiyama A., Kawakami T., Sugano S., Nagahari K., Isogai T., Insogai T., Isogai T., Insogai T., Insoga
                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nollo sapreus viunnai,.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..
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0
                                                                                     Length 143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 30.0%; Score 6; DB 2; Length 146; Best Local Similarity 100.0%; Pred. No. 57; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                             Indels
1
143 143
143 AA; 15957 MW; E56ABB6F1F30879F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7186; BAC86876.1; -.
146 AA; 16128 MW; C41FD600CF9E8FD1 CRC64;
                                                                                                                                                                                                                                                                                                                                       05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein FLJ45251.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
CDNA FLA45251 fis, clone BRHIP2009177.
                                                                                     DB 2;
                                                                              30.0%; Score 6; DB 2; 100.0%; Pred. No. 56; Live 0; Mismatches
                                                                                                                                                                                                                                                                                                               146 AA
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                                                                          Query Match
Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                         PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Hippocampus;
                                                                                                                                                                                                     11 LGWRWL 16
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BAC86876;
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Q6ZSS0
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"Clade analysis and surface antigen polymorphism of hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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NCBI_TaxID=10407;
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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                                                                                                                                   Indels
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EMBL; AK127186; BAC86876.1; -. SEQUENCE 146 AA; 16128 MW; C41FD600CF9E8FD1 CRC64;
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159 AA; 17194 MW; BD6C9235A3E07233 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                 05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
                                                                               30.0%; Score 6; DB 2;
100.0%; Pred. No. 57;
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              American genotypes.",
J. Med. Virol. 72:377-384 (2004).
EMBL, AY311353; AAQ86848.1; -.
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
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Med. Virol. 72:377-384(2004).

EMBL, AY311353; AAQ86848.1;

NON_TER 159 159
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Hepatitis B virus.
                                                                                                                             6; Conservative
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185 1
185 AA;
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                                                                                                                                          Smr domain protein.
WLRRYG 17
                    54 WLRRYG 59
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Pubmed=15064399; DOI =00. 1073/pnas.0307639101;

A Seshadri R., Myers G.S.A., Tettelin H., Eisen J.A., Heidelberg J.F.,

A Bodson R.J., Davidsen T.M., DeBoy R.T., Fouts D.E., Haft D.H.,

Selengut J., Ren O., Brinker L.M., Madupu R., Kolonay J.F.,

A Gebregoergis E., Ger K., Tsegave G., Malek J.A., Ayodeji B.,

Shatsman S., McLeod M.P., Smang O., Sodergren E., Baca E.,

A withouther pinchete genomes ".; Knang O., Sodergren E., Baca E.,

Mith other spirochete genomes ".; Fraser C.M., Paulsen I.T.;

Mith other spirochete genomes ".; Rulsen Treponema denticola with other spirochete genomes" in pathogen Treponema denticola proc. Natl. Acad. Sci. U.S.A. 101:5646-5651(2004).

REMBL, ABE017259; AASI1772.1; "."
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Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.
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Bacteria, Proteobacteria, Gammaproteobacteria, Pseudomonadales,
Pseudomonadaceae, Pseudomonas.
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                                                                                                                                                                                                                                                                                               30.0%; Score 6; DB 2; Length 162; 100.0%; Pred. No. 62; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                         Kim J.J., Lawrence J.E., Kluepfel D.A.;
Submitted (OCT-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY163858; AAP30022.1; -.
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       170 AA; 19438 MW; 9A29EB2307D04319 CRC64;
                                                                                                                                                                                                                                                                          162 AA; 19205 MW; 6AD69FCF5A3689DA CRC64;
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Last annotation update)
                                                                                                       01-JUN-2003 (TrEMBLrel. 24, Created)
01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30.0%; Score 6; DB 2; ilarity 100.0%; Pred. No. 64; Conservative 0; Mismatches
                                                                                     162 AA.
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STRAIN=ATCC 35405 / DSM 14222;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OrderedLocusNames=TDE1254;
                                                                                                                                                                                                                                                                                                               Local Similarity 100.
nes 6; Conservative
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SMART; SM00463; SMR; 1.
                                                                                      PRELIMINARY;
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Best Local Similarity
6; Conserve
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                            154 SSWALG 159
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      SSWALG 9
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Matches
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Q73NA1
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MEĎLINE-22196947; PubMed-12209321;
Quintero A., Martinez D., Alarcon De Noya B., Costagliola A.,
Urbina L., Gonzalez N., Liprandi F., Castro De Guerra D., Pujol F.H.;
"Molecular epidemiology of hepatitis B virus in Afro-Venezuelan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                  Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema
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NCBI_TaxID=10407;
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-UNN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
                                                   Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
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6; Conservative 0; Mismatches
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Arch. Virol. 147.1829-1836 (2002).
Arch. Virol. 147.1829-1836 (2002).
GO, GO:0016032; Priviral life cycle; IEA.
InterPro; IRR000349; Hepvir_surfAg.
Pf00695; vMSA; 1.
                                     Created)
PRT;
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   PRELIMINARY;
                   AAS11772;
02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
04-MAY-2004 (TrEMBLrel.
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                                            Query Match
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Q9DQV3
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Quintero A., Martinez D., Alarcon De Noya B., Costagliola A.,
Urbina L., Gonzalez N., Liprandi F., Castro De Guerra D., Pujol F.H.;
"Molecular epidemiology of hepatitis B virus in Afro-Venezuelan
                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=22196947; PubMed=12209321;
Quintero A., Martinez D., Alarcon De Noya B., Costagliola A.,
Urbina L., Gonzalez N., Liprandi F., Castro De Guerra D., Pujol F.H.;
"Molecular epidemiology of hepatitis B virus in Afro-Venezuelan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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NCBL_TaxID=10407;
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        Length 185;
                                                          Indels
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20716 MW; EF100280E26BA81A CRC64;
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20672 MW; 4A1275F2976BAD19 CRC64;
                                                                                                                                                                                                                                                                                                                                           01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-UUN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
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01-UND-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
Hepatitis B virus.
        DB 2;
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30.0%; Score 6; DB 2
100.0%; Pred. No. 69;
iive 0; Mismatches
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EMBL; AF479494; AAN28697.1; -.
05. GO:00106032; P:Viral life cycle; IEA.

InterPro; IPR000349; Hepvir_surfAg.

Pfam; PF00695; vMSA; 1.
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                                                          6; Conservative
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185 1
185 AA;
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Matches 6; Conserv
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"Hepatitis delta virus genotypes I and III circulate associated with
"Hepatitis B virus genotype F In Venezuela.";
J. Med. Virol. 64:356-359(2001).
EMBL. AF251386; AAG44621.1;
EMBL. AF251386; AAG44621.1;
InterPro; IPR000349; HepVir SurfAg.
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Illarramendi X., Guevara M.E., Ludert J.E., Blitz L., Liprandi F.,
Pujol F.H.,
Hepatitis delta virus genotypes I and III circulate associated with
hepatitis B virus genotype F In Venezuela.";
J. Med. Virol. 64:356-359(2001).
BMBL, AF251385; AAG4620.1;
InterPro, IPR000349; Hepvir_surfAg.
Pfam; PF00695; VMSA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=21316770; PubMed=11424126;
Quintero A., Uzcategui N., Loureiro C.L., Villegas L.,
Illarramendi X., Guevara M.E., Ludert J.E., Blitz L., Liprandi F.,
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Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus
          Length 185;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-UUN-2003 (TrEMBLrel. 24, Last annotation update)
Surface antigen (Fragment).
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
       DB 2;
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100.0%; Pred. No. 69;
tive 0; Mismatches
                                   69
30.0%; Score 6; DB 2
100.0%; Pred. No. 69;
cive 0; Mismatches
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                         Local Similarity 100.
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0; Gaps Query Match 30.0%; Score 6; DB 2; Length 185; Best Local Similarity 100.0%; Pred. No. 69; Matches 6; Conservative 0; Mismatches 0; Indels

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4 SSWALG 9 |||||| 137 SSWALG 142

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Search completed: October 26, 2004, 07:22:22 Job time : 93.5 secs

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Compugen Ltd.  Search time 85 Seconds (without alignments)	Million cell updates/sec		of the result being printed, score distribution.  Description  Aaol8008 Intracell Aar89346 Cdk2-inte Aav32121 Interacti Aay49340 Cdk2 inte Aay63240 Intracell Aab682724 Pseudomon Aaol8007 Intracell Aab618007 Intracell Aab618928 Cancer-re Aab51899 Human sec Aab51899 Human sec Aab51899 Human sec Abb11899 Human sec Abb11898 Propionib Abm55080 Propionib Abm64638 Propionib Abm85681 Propionib Abm39268 Predomon Abb11084 Human GPC Abo78072 Pseudomon Abg19443 Novel hum Adm26747 Hyperther
GenCore version 5.1.  Copyright (c) 1993 - 2004 Comp  OM protein - protein search, using sw model  Run on: October 26, 2004, 06:46:31; Sear	Title: Perfect score: 20 Sequence: 1 QVWSSWALGWRWLRRYGWGM 20 Scoring table: 01.IGO Gapop 60.0 , Gapext 60.0 Searched: 2002273 seqs, 358729299 residues	num DB seq length: 0 mum DB seq length: 200000000  -processing: Listing first 100 summans est that is geneseqp1980s:* 1: geneseqp2000s:* 2: geneseqp2000s:* 3: geneseqp2001s:* 5: geneseqp2001s:* 6: geneseqp2001s:* 7: geneseqp2003s:* 8: geneseqp2003s:*	Pred. No. is the number of results predicted the score greater than or equal to the score of the and is derived by analysis of the Lotal score of the Match.  No. Score Match Length DB ID  2 10 100 20 5 AAN9346 3 13 65.0 20 2 AAN93121 4 13 65.0 20 3 AAN9340 5 8 40.0 480 7 ABN9440 6 7 35.0 17 5 AAN93340 7 35.0 66 4 AAN952121 11 6 30.0 55 3 AAB33896 9 6 30.0 55 3 AAB33896 10 6 30.0 66 4 ABB15989 11 6 30.0 66 4 ABB15989 11 6 30.0 97 6 ABN55080 12 6 30.0 112 5 ABN55080 14 6 30.0 112 5 ABN5437 15 6 30.0 114 6 ABN32068 16 30.0 114 6 ABN32068 17 ABN32068 18 6 30.0 114 6 ABN32068 18 6 30.0 114 6 ABN32068 19 6 30.0 138 7 ABN32068 20 6 30.0 138 7 ABN32068 21 6 30.0 138 7 ABN32063 22 6 30.0 171 7 ABO78072 23 6 30.0 277 7 ABN56147

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Add18295 P
Aav14653 H
Add18291 H
Add18991 F
Aav14659 H
Adv1450 H
Adv18283 H
Adv18283 H
Adv18283 H
Adv182936 H
Adv182936 H
Adv18295 H
Adv18291 H
Adv1831 H

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AAR89340-R89351 are Cdk-2 interacting peptides identified by an interaction trap assay using conformationally-constrained proteins. The assay comprises providing a host cell (esp. a yeast cell) which contains: the protein gene operably linked to a DNA-binding protein recognition site, pref. LexA, (ii) a 1st fusion gene expressing cdk-2 covalently bonded to a LexA binding protein, and (iii) a 2nd fusion gene expressing a conformationally-constrained (pref. with thioredoxin) protein (either expected to interact with Cdk-2 or a random peptide) covalently bonded to a gene-activating and measuring expression of the reporter gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            as a measure of interaction between Cdk2 and the conformationally constrained protein. The same system may be applied to find proteins that interact with any other protein of interest (e.g. Ras) or proteins having agonist or antagonist activity on such interactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Interaction trap systems using conformationally-constrained proteins - useful for detection of protein interactions and for identification and isolation of interacting proteins.
                                                                                                                                            Cdk2; cyclin dependent kinase 2; assay; identifying; isolating; cell cycle; interaction; antagonist; conformationally-constrained; agonist; interaction trap; thioredoxin; LexA; two-hybrid system.
                                                                                                       Cdk2-interacting peptide isolated using interaction trap assay.
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Pred. No. 9.9e-07;
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100.0%; Pre
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                                                                                                                                                                                                                       Synthetic.
                                AAR89346;
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         Abo00234 Novel hum
Abo11566 Human sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Process for specifically modulating the properties of an intracellular target molecule used for the treatment of various disorders.
                                                                                                                                                                                                                                                                           Intracellular target molecule property modulation method aptamer 10M.
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                                                                              ALIGNMENTS
   AB000234
AB011566
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(MASS-) MASSACHUSETTS GEN HOSPITAL.
(MOLE-) MOLECULAR SCI INST.
                                                                                                                                                                 AA018008 standard; peptide; 20 AA.
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     99
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   1089
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                                                                                                                             RESULT 1
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                                                                                                                                                  AA018008
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0;

Gaps

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Length 20; Indels Protein interaction; interaction trap; fusion protein; mimetic; therapeutic; detection; reporter gene.

Synthetic.

AAR89346 standard; peptide; 20 AA.

RESULT 2 AAR89346 ID AAR8

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Similarity
                                                                                                                                                                                                                                                                        Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rubenfield MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-FEB-1998;
27-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L8-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JS6551795-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-APR-2003.
                                                                                                                                                                                                                                                 polypeptide
                                                               Mccoy JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO82724;
                                                                                                                                                                                                                                                                                              Query Match
Best Local 8
                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 5
                                                                                                                                                                                                                                                                                                                                                                                                              ABO82724
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                                                                                                                                                                                                                                                                                                                                                                g
                                                                                                                                                                                                                                             Aptamers AAW32116-W32132 have been isolated from a peptide library and are used in a novel interaction trap method for detecting protein interactions and isolating novel proteins. The method involves a host call containing a reporter gene operably linked to a DNA-binding-protein (DBP) recognition site, a fusion gene capable of expressing a fusion protein which is able to specifically bind to the DBP recognition site and a second fusion gene which expresses a second fusion protein which is conformationally constrained and bonded to a gene activating modety. Measuring expression of the reporter gene gives a measure of the interaction between Pl and P2. This method can be used to identify agonists or antagonists for use as therapeutic molecules or for the detect an interaction protein in a population of proteins or to identify a candidate interactor. Using conformationally constrained proteins can provide for tertiary structural analysis and can also protect proteins from callular degradation and/or increase the protein's solubility, and/or otherwise alter the capacity of the candidate interactor to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                              New trap system for detecting protein interactions - comprises a reporter gene linked to a DNA-binding-protein recognition site and fusion proteins to test for interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces, mammalian, fusion protein, interactor peptide, conformation-constraining protein, DNA binding moiety, Cdk2, gene activating moiety, protein interaction, gene purification.
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                                                                                                                                   Xu CW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY49340 standard; peptide; 20 AA.
                                                                                                                                                                                                                            Claim 73; Page 52; 89pp; English.
                                                                                                                                   Mccoy JM, Jessen TH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cdk2 interacting peptide i5-4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-00504538,
                                                     97WO-US005793
                                                                            96US-00630052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                  GEHO ) GEN HOSPITAL CORP
                                                                                                              (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 WALGWRWLRRYGW 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WALGWRWLRRYGW 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                         WPI; 1997-512733/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-1995;
                                                     09-APR-1997;
                                                                            09-APR-1996;
         WO9738127-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13;
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                                16-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                   Brent R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY49340,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cells comprising recombinant DNA molecules encoding fusion proteins, each consisting of a candidate interactor peptide, a conformation-constraining protein and a DNA binding molety and/or gene activating molety. The cells are useful for detecting protein interactions. The cells may also be used in a method for identifying and purifying genes encoding a wide range of useful proteins based on their physical interaction with a second
                                                                                                                                                                                                                                                                                                                     Population of Saccharomyces and/or mammalian cells comprising recombinant DNA encoding fusion proteins, useful for detecting protein interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a population of Saccharomyces and/or mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65.0%; Score 13; DB 3; Length 20; 100.0%; Pred. No. 9.9e-07; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 31470; 455pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nolling J, Deloughery C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pseudomonas aeruginosa polypeptide #14899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO82724 standard; protein; 480 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 3B; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENO-) GENOME THERAPEUTICS CORP.
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                                                                                                                                                                           Jessen TH, Brent
20-JUL-1994; 94US-00278082
                                                               (GEHO ) GEN HOSPITAL CORP. (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 WALGWRWLRRYGW 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WALGWRWLRRYGW 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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N-PSDB; ABD16295.
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            prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa mucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa durgs, as templates for recombinant production of P. aeruginosa-derived peptides or polypeptides, as target components for diagnosis and/or treatment of P. aeruginosa-causaed infection, and in detection of P. aeruginosa sequences or other sequences of Pseudomonas species using biochip technology. Sequences ABO67826-ABO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a process for specifically modulating the properties of an intracellular target molecule T, and/or of a cellular component C which interacts directly or indirectly in a cell with the target. The process involves the introduction into the cell of chimeric molecule known as a targeted effector, comprising a recognition moiety capable of recognising T and an effector moiety. The chimeric protein or nucleic acid can be used in the preparation of a medicament for the treatment of microbial infections, immunological disorders,
  pathological conditions, as molecular targets for diagnostics,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antimicrobial; immunomodulatory; nootropic; neuroprotective; metabolic; neuroleptic; cytostatic; cardiant; infection; immunological disorder; neurological disorder; metabolic disorder; psychiatric disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Process for specifically modulating the properties of an intracellular target molecule used for the treatment of various disorders.
                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Intracellular target molecule property modulation method aptamer 10.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Intracellular target; cellular component; property modulation;
                                                                                                                                                                                                                                                                                                                                     Length 480;
                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                   DB 7;
                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            myopathy; cancer; cardiovascular disorder
                                                                                                                                                                                                                                                                                                                                 40.0%; Score 8; I
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MASS-) MASSACHUSETTS GEN HOSPITAL.
(MOLE-) MOLECULAR SCI INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA018007 standard; peptide; 17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-NOV-2000; 2000EP-00403156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-2000; 2000EP-00403156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cohen BA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CNRS ) CENT NAT RECH SCI.
                                                                                                                                                                                                                                                                                                                                                                        8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Fig 1; 33pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      262
                                                                                                                                                                                                                                                                                                                                                                                                                 8 LGWRWLRR 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-418829/45.
                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Brent R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    255 LGWRWLRR
                                                                                                                                                                                                                                                                                              Sequence 480 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA018007;
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                                                                                                                                                                                                                                                                                                                                                                          Matches
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The polynucleotide sequences given in AAF32699 to AAF32747 encode the human secreted proteins given in AAB64422 to AAB64470. AAB64471 to AAB64548 represent human secreted polypeptide sequences and proteins (AAB64548 represent human secreted polypeptide sequences and proteins homologous to them, which are given in the exemplification of the present invention. Human secreted proteins have activities based on the tissues and calls the genes are expressed in Examples of activities include: antiarthritic; immunosuppressive; antirheumatic; antipoliferative; cytostatic; cardiant; vasotropic; cerebroprotective; notropic; neuroprotective; antibacterial; vincide; fungicide; and polypeptides can be used to prevent, treat or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are also used in diagnosing a pathological condition or susceptibility to a pathological condition. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative
                                                                                                                                                                                  0;
neurological disorders, metabolic disorders, psychiatric disorders, wopathles, genetic disorders, candicovascular disorders and dental disorders. The present sequence is a known anti-Cdk2 aptamer used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, secreted protein, diagnosis; immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; neoplasm; cardiovascular disorder; arrest; cerebrovascular disorder; cerebroma ischemenia; infection; angiogenesis; nervous system disorder; Alzheimer's disease; skin aging; ocular disorder; conneal infection; wound healing; food additive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated nucleic acid molecule encoding a human secreted protein is used in preventing, treating or ameliorating a medical condition.
                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted protein sequence encoded by gene 19 SEQ ID NO:78.
                                                                                                                                                                                  0;
                                                                                                                                                                                  0; Indels
                                                                                                                                                     100.0%; Pred. nc.
                                                                                                                                               Score 7; I
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 520-521; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                     AAB64440 standard; protein; 66 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                               35.0%;
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                                                                                                                                                                                  7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-025337/03.
                                                                                                                                                                                                                        6 WALGWRW 12
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                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; AAF32717
                                                                                                    Sequence 17 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preservative.
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                                                                                                                                                                                                                                                                                                                                                                                                         AAB64440;
                                                                                                                                                                                  Matches
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Sequence 18 AA;

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disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel method of selecting target and target binder pairs comprising mixing in a reaction vessel phage expressing biological targets and phage expressing target binders, each having distinguishable selection markers and selecting target and target binder pairs based on the selection markers. The molecules of the invention demonstrate cytostatic activity whilst the method may be useful for selecting target and target binder pairs for preparing a composition for treating cancer. Furthermore, the method may be utilised during gene therapy procedures. The current sequence is that of the cancer-related DGI-2-binder peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Selecting target and target binder pairs for preparing a composition for treating cancer by mixing in a reaction vessel phage expressing biological targets and phage expressing target binders.
                                                                                                                                     culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities, AAR32690 to AAR32698 and AAB64421 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; cancer; gene therapy; DGI-2; DGI-5; DGI-7; DGI-9; Hras; leptin; VEGF; vascular endothelial growth factor receptor; VEGF-R1; VEGF-R2; VEGF-R3; FLT1; FMS-related tyrosine kinase 1; FLK1; KDR; kinase insert domain protein receptor; EGFR; epidermal growth factor;
                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                   Length 66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cancer-related DGI-2-binder peptide - SEQ ID 115.
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                                                                                                                                                                                                                                                                                                   DB 4;
                                                                                                                                                                                                                                                                                               35.0%; Score 7; DB 4
100.0%; Pred. No. 4.9
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FGFR1; fibroblast growth factor; Tie-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC99282 standard; peptide; 18 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-OCT-2001; 2001US-0345471P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Brissette R, 8
Goldstein N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100...
7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                               3 WSSWALG 9
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                                                                                                                                                                                                                                                              Sequence 66 AA;
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Prendergast J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
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                                                                                                                                                                                                                         nvention
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The invention relates to the isolation of genes AAC59335-C59382 encoding the human secreted proteins AAB3844-B33891. The sequence is a search result from a BLASTX homology search. The genes and proteins are useful for preventing, ameliotrating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer, and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal cract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, theumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epileppy; and (f) infectious diseases such as viral, bacterial, fungal and parasitic infections
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                                                                                                                                                                                                                                                                                                                                               Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiuleer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder; neurological disease; infection; human; secreted protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted proteins and the nucleic acids that encode them, useful in gene therapy protocols and recombinant nucleic acid based procedures.
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                                                                                                                                                                                                                                                                                                              Human secreted protein BLAST search protein SEQ ID NO: 111.
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                                               0; Indels
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           Length 18;
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            DB 7;
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              Score 6; DB 7
Pred. No. 19;
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                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 377; 410pp; English.
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                                                                                                                                                                                                         AAB33896 standard; protein; 55 AA.
30.0%; SCC_
100.0%; Pre
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                                                 6; Conservative
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Best Local Similarity
Matches 6; Conserv
                           Local Similarity
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                                                                                  9 MSSMAO
                                                                                                                   4 QVWSSW 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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            Query Match
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                                                 Matches
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05-SEP-2000; 2000US-0229513P.
06-SEP-2000; 2000US-0231243P.
08-SEP-2000; 2000US-0231244P.
08-SEP-2000; 2000US-0231244P.
08-SEP-2000; 2000US-0231244P.
08-SEP-2000; 2000US-0231244P.
08-SEP-2000; 2000US-0231244P.
08-SEP-2000; 2000US-0231444P.
08-SEP-2000; 2000US-023144P.
08-SEP-2000; 2000US-0231444P.
08-NOV-2000; 2000US-0241869P.
08-NOV-2000; 2000US-0246477P.
08-NOV-2000; 2000US-0246477P.
08-NOV-2000; 2000US-0246522P.
08-NOV-2000; 2000US-024652P.
08-
       Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antiatickling; antianaemic; antiarthritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder; neurological disease; infection; nephrotropic; gene therapy; vaccine.
                                                                                                                                                                                                                                                                 Human nervous system related polypeptide SEQ ID NO 4646
                                                                                                                                                   ABB15989 standard; protein; 66 AA
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2000US-0186528P.
2000US-0186528P.
2000US-0186530P.
2000US-0186874P.
2000US-0198874P.
2000US-0198874P.
2000US-01988874P.
2000US-0218486P.
2000US-021487P.
2000US-021487P.
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2000US-021487P.
2000US-021487P.
2000US-021487P.
2000US-021487P.
2000US-021487P.
2000US-022584P.
2000US-022584P.
2000US-022584P.
2000US-022588P.
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2000US-0229343P.
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2000US-0229509P
                                                                                                                                                                                                                            (first entry)
                                                        SSWALG 12
                       σ
                       SSWALG
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14-AUG-2000; 2
14-AUG-2000; 2
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18-AUG-2000; 2
22-AUG-2000; 2
22-AUG-2000; 2
22-AUG-2000; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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Annexin binding epitope, hepatitis B virus; hepatitis D virus; influenza; benzodiazepine; binding inhibitor; cytomegalovirus; viral infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a fragment of the hepatitis B virus surface antigen F amino acid sequence. The antigen sequence is used to determine the best sequence to use as a source of peptides for the composition of the invention. The invention relates to benzodiazepine derivatives
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New benzodiazepine and benzothiazepine derivatives, useful for treating or preventing viral infection, contain peptides that include an annexinbinding epitope.
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                      Hepatitis B virus surface antigen F amino acid sequence.
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100.0%; Pred. No. 58;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                               99WO-EP006231.
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                            Hepatitis B virus.
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                                                                                                                                                  benzothiazepine
                                                                                                                                                                                                                                                                         WO200012547-A2
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$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac{1}{2}$\frac
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acids encoding 3224 human nervous system antigen polypeptides, useful for preventing, diagnosing and/or treating nervous system cancers and metastases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ruben SM;
                                                                                                               2000US - 0249264P.
2000US - 0249265P.
2000US - 0249297P.
2000US - 024929P.
2000US - 02500P.
200US - 0250191P.
                                                                                                                                                                                                                                                                                                                                     2000US-0251030P.
2000US-0251988P.
                                                                                                                                                                                                                                                                                                                                                                                               2000US-0256719P.
2000US-0251479P.
2000US-0251856P.
                            2000US-0249218P.
                                                                                         000US-0249245P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0251990P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-0251869P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0251989P
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Best Local Similarity Ivv.
6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-541565/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SWALGW 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; ABA12315
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                                                                                                                                                                               17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                  05-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JAN-2001;
                                                                                                                  17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                            -DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-DEC-2000;
                                                                                                                                                                                                                                               17-NOV-2000;
                                                                                                                                                                                                                                                                            01-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                     05-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-DEC-2000;
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20-JUN-2000
                                                                                                                                                                                                                                                                                                                                         2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ŋ
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                                                                                            -7-NOV-7
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Gaps

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0; Indels

RESULT 11

ð a AAYB5124 ID AAYE XX AC AAYE XX DT 06-7 DT 20-5

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Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophhalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies polypeptides may be used as antigens in the production of antibodies of specific for P. acnes proteins. These antibodies may also be used to downregulate expression and activity of P. acnes polypeptides and claiming P. acnes presence, for example, by diagnostic agents for determining P. acnes presence, for example, by claim patent did not form part of the printed specification, but was contacting in the printed specification, but was contacting the polyment of the printed specification, but was contacting and activity of P. acnes presence data for child and discontacting assay (ELISA). Note: The sequence data for child activity of the printed specification, but was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; SEQ ID NO 19756; 1069pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         L'maisonneuve J, Zhang Y, Jen S,
                                                                                                                                                                                                                    21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                              20-APR-2001; 2001WO-US012865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treating acne vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-616774/71.
                                                                                                                                                                                                                                                                                                                                                           (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; AAS59592
          WO200181581-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                 Skeiky YAW,
                                                                            01-NOV-2001
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          g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to isolated polymucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain recardion (PCN) primers, oligomers, and for chomosome and gene mapping, and in recombinant production of (II). The polymucleotides are also used in diagnostics as expressed sequence tags for identifying expressed entitly for its useful in gene therapy techniques to restore normal cativity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of supplement. (II) and its binding partners are useful for treating disorders of supplement. (II) and (II) are useful for treating disorders of polypeptide and polymucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations and to produce other types of data and products dependent on DNA and amino acid sequences. Abg00010-Abg30377 represent novel human diagnostic amino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in the form of the printed specification, but was obtained in the contract of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 20; SEQ ID NO 43911; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                           30-MAR-2001; 2001WO-US008631
                                                                                                                                                                                                                                                                                                                                                                                            Tang YT;
                                                                                                                                                                                                                31-MAR-2000; 2000US-00540217
                                                                                                                                                                                                                                                 2000US-00649167
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                                                                                                                                                                                                                                                                                                                                                                                     Drmanac RT, Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-639362/73
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                   (HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAS77739
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WO200175067-A2,
                                                                                                                                                                                                                                                 23-AUG-2000;
                                                                 11-0CT-2001
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immunogenic

Bhatia A;

Wang SS,

Mitcham JL, Walls

Persing DH,

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0
                                                                         Gaps
                                                                                                                                                                                                                                                        Propionibacterium acnes predicted ORF-encoded polypeptide #19756.
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                                                                                                                                                                                                                                                                              Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                        0; Indels
                                                 Length 97;
                                                 4,
                                              Score 6; DB 4;
Pred. No. 74;
0; Mismatches
                                                                                                                                                                                                                                                                                            immunostimulant; immune response; vaccine.
ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                 ABM55080 standard; protein; 97 AA.
                                                      100.0%; Pr
                                                 30.0%;
                                                                                                                                                                                                                                  20-OCT-2003 (first entry)
                                                                        6; Conservative
                                                                                                                                                                                                                                                                                                                    Propionibacterium acnes
                                              Query Match
Best Local Similarity
                                                                                                                      68 WSSWAL 73
                                                                                               3 WSSWAL 8
                                                                                                                                                                                                                                                                                                                                          WO2003033515-A1.
                        Sequence 97 AA;
                                                                                                                                                                                                         ABM55080;
                                                                                                                                                       RESULT 14
ABM55080
ID ABM550
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SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endopthalmitis; bone; joint; central nervous system; EliSA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.

Propionibacterium acnes

Propionibacterium acnes immunogenic protein #19457.

(first entry)

13-FEB-2002 AAU58561;

AAU58561 standard; protein; 97 AA.

AAU58561

g à

Maisonneuve JL; Jones R, Carter D;

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Acne vulgaris, antiseborrhoeic, dermatological; antibacterial; immunostimulant; immune response; vaccine; immunogenic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Propionibacterium acnes immunogenic polypeptide #29314.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM64638 standard; protein; 98 AA.
                                                     Mitcham JL, Skeiky YAW, Persi
Zhang Y, Wang S, Jen S, Lode
Barth B, Vallieve-Douglass J;
             11-OCT-2002; 2002WO-US032727.
                           15-OCT-2001; 2001US-00978825.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                    Query Match
Pest Local Similarity 100...
6: Conservative
                                        (CORI-) CORIXA CORP
                                                                                 WPI; 2003-381789/36
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                                                                                                                                                                                                                                                                                                                                                                                        3 WSSWAL 8
                                                                                         N-PSDB; ACF64521
                                                                                                                                                                                                                                                                                                                                               Sequence 97 AA;
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24-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                      68
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM64638;
                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
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The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polymetric fusion proteins comprising a polypeptide of the invention, an immunogenic specific for a P acnes of the invention, fusion proteins comprising a polypeptide of the invention; and an isolated T cell population comprising T cells prepared via this method; a vaccine composition (comprising P. acnes polypeptides, this method; a vaccine composition (comprising P. acnes polypeptides, polymerications, or antipolies, fusion proteins, T cell populations, or antipolies, antibodies, fusion proteins, T cell populations or antipol-persenting or determining the presence or absence of P. acnes in a patient; and a method for inhibiting the development of P. acnes in a patient; and a method for inhibiting the development of P. acnes in a proteins, T cell populations or antipol-presenting or treating acnes polypeptides an immune response specific for a P. acnes to polypeptides are useful for diagnosing, preventing or treating acnes protein. The polymucleotides can also be used as probes or primers for nucleic acid hybridisation. The vaccine composition is useful for the immune response against P. acnes, or for treating acnes and the kit is useful for performing a diagnostic assay. The present contain an immunogenic region. Note The sequence data for this patent did not form part of the printed specification, but was the contain an immunogenic region of the sequence data for this patent did not form part of the printed specification, but was contained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New Propionibacterium acnes polypeptides and polynucleotides encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                                                                                                                                                 Bhatia A,
Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30.0%; Score 6; DB 6
100.0%; Pred. No. 74;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 7; SEQ ID NO 29314; 1481pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           obtained in electronic format directly f:
ftp.wipo.int/pub/published pct sequences
                                                                                                                                                                                                                                                                                                                                                                                    Persing DH,
                                                                                                                                                                                                                                                                                                                                                                                                                      Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABP64437 standard; protein; 112 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                  Zhang Y, Wang S, Jen S, Lod
Barth B, Vallieve-Douglass J;
                                                                                                                                                                                        11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                     L5-OCT-2001; 2001US-00978825
                                                                                                                                                                                                                                                                                                                                                                                 Mitcham JL, Skeiky YAW,
Zhang Y, Wang S, Jen S,
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Propionibacterium acnes.
                                                                                                                                                                                                                                                                                                                    (CORI-) CORIXA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 WSSWAL 8
                                                            WO2003033515-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 98 AA;
                                                                                                                            24-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynucleotide of the invention; antibodies against polypeptides of the invention; invention; fusion proteins comprising a polypeptide of the invention; a method for stimulating an immune response specific for a P. acnes polypeptide and an isolated T cell population comprising T cells prepared polypeptide and an isolated T cell population comprising T cells prepared to a vaccine composition (comprising P. acnes polypeptides, polynucleotides, antibodies, fusion proteins, T cell populations, or antigen-presenting cells that express the polypeptides, antibodies, fusion proteins, T cell populations or antigen-presenting cells that express the polynucleotides, antibodies, fusion proteins, T cell populations or antigen-presenting cells that express the polypeptides are useful for diagnosing, preventing cells that express the proteins, T cell populations or antigen-presenting cells that express the proteins. T cell populations or antigen-presenting cells that express the proteins, or for stimulating an immune response specific for a P. acnes for mucleic acid hybridistation. The polynucleotides can also be used as probes or primers for remaining an immune response expectition of an immune response expective and the protein and primers for the polynucleotides can also be used as probes or primers for etimulations of an immune compassition its useful for the polynucleotides.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New Propionibacterium acnes polypeptides and polynucleotides encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                        Maisonneuve JL;
Jones R, Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Erom WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                     Bhatia A,
Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; DB 6;
100.0%; Pred. No. 74;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; SEQ ID NO 19756; 1481pp; English.
                                                                                                                                                                                                                                                        Persing DH,
Lodes MJ,
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· 0

Gaps

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0; Indels Length 98;

04-NOV-2002 (first entry)

ABP64437;

Human ORF807

DB 6;

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to novel human ORFX polypeptides and their coding sequences (ABP63631-ABP64681 and ABQ98194-ABG99267). The sequences were discovered in human atherogenic cells, in particular in platelets and human umbilital vein endothelial cells (HUVEC) and are expressed in many other tissues as well. Atherogenic cells are cells which have the potential to develop atherosclerotic plaques. The ORFX polypeptides and nucleic acids are useful for treating or preventing a pathological condition associated with an ORFX-associated discase, each cancer, cardiovascular disease, allergy, autoimmune disease, wound healing, blood for this patent did not form part of the printed specification, but was obtained in electronic format directly from the USPTO web site at sequence.html?DocID=20020082206
                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide designated ORFX are present in human atherogenic cells and are useful to prevent and treat ORFX-associated disorders including cancer, allergy, wound healing or autoimmune, cardiovascular or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectent.
                 Antiinflammatory; gene therapy; human; ORFX; atherogenic; platelet; human umbilical vein endothelial cell; HUVEC; atherosclerotic plaque; cancer; cardiovascular disease; allergy, autoimmune disease; wound healing; blood coagulation disorder; inflammatory disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
Cytostatic; Cardiant; Anti-allergic; Immunosuppressive; Vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                        Law D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Propionibacterium acnes immunogenic protein #3645.
                                                                                                                                                                                                                                                                                                                                                                      Topper JN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 10; SEQ ID NO 1614; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 6; D; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                      Conley PB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU42749 standard; protein; 114 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30.0%;
100.0%;
                                                                                                                                                                                                  30-MAY-2001; 2001US-00867550
                                                                                                                                                                                                                                     30-MAY-2000; 2000US-0208427P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Conservative
                                                                                                                                                                                                                                                                                                                                                                      Leach MD, Mehraban F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammatory disease.
                                                                                                                                                                                                                                                                   LEACH M D.
MEHRABAN F.
CONLEY P B.
TOPPER J N.
LAW D.
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-626554/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              σ
                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; ABQ99000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 112 AA;
                                                                                                                                   US2002082206-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 WLRRYG
                                                                                                   Homo sapiens
                                                                                                                                                                    27-JUN-2002
                                                                                                                                                                                                                                                                                  (MEHR/) N
(CONL/) (TOPP/) 7
(LAWD/) 1
                                                                                                                                                                                                                                                                    (LEAC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SABHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (ELISA). Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Propionibacterium acnes predicted ORF-encoded polypeptide #3944.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bhatia A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Length 114;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wang SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mitcham JL, Wang St
Jen S, Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30.0%; bcc. No. c., 100.0%; Pred. No. c., 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 3944; 1069pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ftp.wipo.int/pub/published pct sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM39268 standard; protein; 114 AA.
                                                                                                                                                                                                                                                                         21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                                                                                       20-APR-2001; 2001WO-US012865.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Skeiky YAW, Persing DH, M
L'maisonneuve J, Zhang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Conservative
Propionibacterium acnes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Propionibacterium acnes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating acne vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-616774/71.
                                                                                                                                                                                                                                                                                                                                                                                                                      (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96 VWSSWA 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAS59518.
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                                                                     WO200181581-A2.
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                                                                                                                                      01-NOV-2001
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Bacterial infection; Pseudomonas aeruginosa infection; antibacterial

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The invention relates to an isolated polynuclectide (ACF64435-ACF64733)

encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides. The invention and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a colynucleotide of the invention; antibodies against polypeptides of the invention; polypeptide of the invention; dual of the invention; and invention and an isolated T cell population comprising T calls prepared to polypeptide and an isolated T cell population comprising T. acnes polypeptides, polymention; antibodies, fusion proteins, T cell populations, or polynucleotides, antibodies, fusion proteins, T cell populations, or antigen-presenting cells that express the polypeptide, antibodies, fusion proteins, T cell populations, or patient, and a method for inhibiting the development of P. acnes in a patient, and a method for inhibiting the development of P. acnes in a proteins. T cell populations or antigen-presenting cells that express the polypeptides are useful for diagnosing, preventing or treating acne polypeptides are useful for diagnosing, preventing or treating acne collypeptides are useful for diagnosing, preventing or treating acnes protein. The polymucleotides can also be used as probes or primers for nucleic acid hybridisation. The vaccine composition is useful for the sequence represents a polypeptide predicted to be encoded by an ORF (open the kit is useful for performing a diagnostic assay. The present cand the kit is useful for the vaccine composition assay. The present cand the kit is useful for performing a diagnostic assay. The present cand the kit is useful for performing a diagnostic assay. The present canding trame) contained within the P. acnes polymented presents a polypeptide predicted to be encoded by an ORF (open reading frame) contained within the P. acnes polymented prediction in electronic format directly the printed specification, but was ob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                               New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                               Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
0
                                                                                                                                                                                                                                             Maisonneuve
Jones R, Ca
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                                                                                                                                                                                                                                             Bhatia A,
Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30.0%; Score 6; DB 6;
100.0%; Pred. No. 84;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 3944; 1481pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pseudomonas aeruginosa polypeptide #16263.
                                                                                                                                                                                                                                          JL, Skeiky YAW, Persing DH,
Wang S, Jen S, Lodes MJ,
Vallieve-Douglass J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO84088 standard; protein; 138 AA.
                                                                                                                                                    15-OCT-2001; 2001US-00978825
                                                                                                          11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Conservative
                                                                                                                                                                                                (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                WPI; 2003-381789/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VWSSWA 101
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                                                                                                                                                                                                                                                                                                                                                          N-PSDB; ACF64447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 114 AA;
                  WO2003033515-A1
                                                                                                                                                                                                                                             Mitcham JL,
                                                              24-APR-2003
                                                                                                                                                                                                                                                               Zhang Y,
Barth B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to Pseudomonas aeruginosa polypeptides and the polymuclectides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a pacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templaces for recombinant production of P. aeruginosa drugs, as templaces for recombinant components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences of pseudomonas species using biochip technology. Sequences ABOS 7826-ABOS 4396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; cytokine; cell proliferation; cell differentiation; growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       haematopoiesis regulation, tissue growth, immunomodulator; activin, inhibin; chemotaxis, chemokinesis, thrombolysis, oncogenesis, proliferation; metastasis, cancer; tumour, haematopoietic disorder, myeloid cell disorder; lymphoid cell disorder; asthma; arthritis; chronic inflammatory condition, proliferative retinopathy; atherosclerosis, coronary heart disease, arterial ischaemia; bone disorder; osteoporosis, vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   specification but was obtained in electronic format from USPTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                            Rubenfield MJ, Nolling J, Deloughery C, Bush D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted protein homologue, SEQ ID NO:1454.
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Pred. No. 98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30.0%; Scor.
100.0%; Pred. No. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 32834; 455pp; English
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                                                                                                                                                                                                                                       (GENO-) GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      seqdata.uspto.gov/sequence.html
                                                                                                                                                                                 98US-0074788P.
98US-0094190P.
                                                                                                                                              99US-00252991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
                                   Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                WPI; 2003-615309/58.
N-PSDB; ABD17659.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 138 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10 WRWLRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49 WRWLRR
                                                                                                                                                                                 18-FEB-1998;
27-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-JAN-2002
                                                                                                                                              .8-FEB-1999;
                                                                                                           22-APR-2003.
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cell culture; drug screening; gene therapy; antiinflammatory; antiathmatic; antiarthritic; haemostatic; antiarteriosclerotic; cytostatic; osteopathic; vasotropic; cardiant; virucide; antibacterial; antifungal; vulnerary; antiulcer.
                                                                                                                                                       Claim 20; Page 144; 1963pp; English.
                                                                                                    Liu C, Drmanac RT;
                                                             05-FEB-2001; 2001WO-US003800
                                                                        03-FEB-2000; 2000US-00496914
27-APR-2000; 2000US-00560875
                                                                                                               2001-457740/49.
                                                                                         (HYSE-) HYSEQ INC.
                                                                                                                     N-PSDB; ABA08328
                                       WO200157188-A2
                            Homo sapiens.
                                                                                                                                            and cancer.
                                                                                                    Tang YT,
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Sequences ABB10981-ABB12330 represent 1350 novel human polypeptides, and caguences AABB10981-ABB12330 represent nuclear caids encoding them. The invention also relates to vectors and recombinant host cells comprising a nucleotide of the invention, methods of producing the novel polypeptides. C antiblodies against the polypeptides, methods of dentifying compounds which bindess against the polypeptides of identifying compounds which polypeptides of the invention have homology to known proteins, thereby giving an insight into their probable biological activities, and hence coptemnial therapeutic applications. The polypeptides of the invention may have various activities; stem cell growth factor activities, of differential therapeutic applications. The polypeptides of the invention may have various activities; stem cell growth factor activity; issue growth activity; immunomodulatory activity; activity, tissue growth activity; is memorated or chemokinetic activities; properties or may be involved in onosogenesis, cancer cell proliferation or metastasis. Chemotocytic activities; receptor or ligand activity; or may be involved in onosogenesis, cancer cell proliferation or metastasis. Coperation or their biological activities; polypeptides and nucleotides of the invention are useful for perventing, treating or ameliorating medical cancers, haemacopoletic disorders (e.g., myeloid or lymphoid cell conditions, e.g., by protein or gene therapy, such conditions include cancers, haemacopoletic disorders (e.g., myeloid or lymphoid cell useful. Polypeptides involved with tissue regeneration and repair is chaemia, bone disorders (e.g., myeloid or lymphoid cell cancers, haemacopoletic disorders (e.g., myeloid or lymphoid cell cancers, haemacopoletic disorders (e.g., costeoprossis), and abnormal vascular growth, Polypeptides involved with tissue so astimate of beating (e.g., of burns, inclsions and ulcers), while those with growth factor activity may be used to promote cell growth. For example, such polypeptides may be used to au
Human proteins and DNA encoding sequences useful for preventing, or ameliorating a medical condition in a mammalian subject e.g. \epsilon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 143 AA;
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triphosphate-binding protein coupled receptor (GPCR). A polymucleotide of the invention may have a use in gene therapy. The polymucleotide and polypeptide are useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor. The protein sequences shown in ADC85549-ADC87617 represent GPCR's of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polynucleotide, useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor.
                                                                                                                                                                                    human; GPCR; guanosine triphosphate-binding protein coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel polynucleotide encoding a guanosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 165;
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                                                                                                                                                                                                                                                                                                                                                                   (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
(ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30.0%; Score 6; DB 7
100.0%; Pred. No. 1.1
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pseudomonas aeruginosa polypeptide #10247.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; SEQ ID NO 1396; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AB078072 standard; protein; 171 AA.
                                                                    ADC86943 standard; protein; 165 AA.
                                                                                                                                                        Human GPCR protein SEQ ID NO:1396.
                                                                                                                                                                                                                                                                                                                                                                                                          Akiyama Y,
                                                                                                                                                                                                                                                                                                                                        18-JUN-2001; 2001JP-00246789.
                                                                                                                                                                                                                                                                                                            18-JUN-2002; 2002EP-00013517
                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-315783/31.
N-PSDB; ADC86942.
94
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                                                                                                                                                                                                                                                                                                                                                                                                             Asai K,
WSSWAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 165 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SWALGW
                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                   EP1270724-A2.
                                                                                                                                                                                                 gene therapy.
                                                                                                                            01-JAN-2004
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                                                                                                                                                                                                                                                                                  02-JAN-2003.
                                                                                                  ADC86943;
68
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                                                                                                                                                                                                                                                                                                                                                                                                            Suwa M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                          RESULT 21
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                                                          ADC86943
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Gaps

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Length 143;

30.0%; Score 6; DB 4; Length 143; llarity 100.0%; Pred. No. 1e+02; Conservative 0; Mismatches 0; Indels

Local Similarity Les 6; Conserv

Query Match Best Loca Matches

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The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (III). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations and to produce other types of data and products dependent on DNA and amino acid sequences. Abgological activity and activity and activity and activity and activity and produces the polymeration of mutations are sequences. Abgological activity and and an analysing and to produce other types of data and products dependent on DNA and and an activity and activity and activity and activity and produces the products dependent on DNA and and an activity and activity and activity and products dependent on DNA and and activity and activity and activity and activity and activity and products dependent on DNA and and activity and activity and activity and products dependent on DNA and activity and activity and activity and products dependent on DNA and activity and activity and products dependent on DNA and activity and activity and activity and activity and products dependent on DNA and and activity and activity and activity and activity and activity and activity and products dependent on DNA and activity and activity and activity and activity and activity and activity and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                       New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 201; . 1.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hyperthermophile; protein stability enhancement;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; DB 4
100.0%; Pred. No. 1.3
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 20; SEQ ID NO 49802; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM26747 standard; protein; 277 AA.
                                                                                                                                                   Tang YT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-MAR-2002; 2002US-0361742P.
14-MAY-2002; 2002US-0380423P.
31-MAR-2000; 2000US-00540217.
23-AUG-2000; 2000US-00649167.
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Best Local Similarity luv...
Si Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100 WSSWAL 105
                                                                                                                                                   Drmanac RT, Liu C,
                                                                                                                                                                                                         WPI; 2001-639362/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 WSSWAL 8
                                                                                          (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 201 AA;
                                                                                                                                                                                                                                    N-PSDB; AAS83630
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                                                                                                                                                                                                                                                                                                                                                                                        biodiversity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant production of P. aeruginosa-derived peptides or polypeptides, as target components for diagnosis and/or treatment of P. aeruginosa-caused components for diagnosis and/or treatment of P. aeruginosa-caused of Pseudomonas species using biochip technology. Sequences ABOS4826 represent P. aeruginosa polypeptides of the invention. Note: The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at sequance.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, chromosome mapping, gene mapping, gene therapy, forensic, food supplement, medical imaging, diagnostic, genetic disorder.
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                                                                                                                                                                                                                                                                                                                                                                                        Bush D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 26818; 455pp; English
                                                                                                                                                                                                                                                                                                                                                                                        Deloughery C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human diagnostic protein #19434.
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                                                                                                                                                                                                                                                                                                                              (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001; 2001WO-US008631.
                                                                                                                                                                                 99US-00252991.
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                                                                                                                                                                                                                                          98US-0074788P
                                                                                                                                                                                                                                                                                                                                                                                        Rubenfield MJ, Nolling J,
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Best Local Similarity الاس.
نام 6; Conservative
            Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-615309/58.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 171 AA;
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                                                                                                                                                                                 18-FEB-1999;
                                                                                                                                                                                                                                                                     27-JUL-1998;
                                                                                                                                                                                                                                          18-FEB-1998;
                                                                  US6551795-B1
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                                                                                                                              22-APR-2003
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therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant production of P. aeruginosa drugs, as templates for recombinant components for diagnosis and/or treatment of P. aeruginosa caused infection, and in detection of P. aeruginosa sequences or other sequences of pseudomonas species using biochip technology. Sequences ABO67826-8BO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence is the woodchuck hepatitis virus surface antigen PreS2-S region. This sequence was used to create a recombinant antigen library. The library comprises recombinant nucleic acids encoding antigenic polypeptides and is produced by recombination of at least two forms of nucleic acid, differing by at least two nucleotides, encoding a disease-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Recombinant multivalent antigenic polypeptide produced by recombining nucleic acid sequences and screening, used in vaccines against e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PreS2-S; recombinant antigen library; disease-related antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          multivalent antigenic polypeptide production; infection; allergen; asthma; autoimmune disease; rheumatoid arthritis; diabetes; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      multiple sclerosis; inflammatory condition; cancer; contraception; immune response; hepatitis b surface antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Woodchuck hepatitis b virus HBsAg PreS2-S region protein sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                               Length 277;
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                                                                                                                                                                                                                                                                                                                                                                                                            DB 7; Ler
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                               Score 6; I
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                           30.0%; Scor
100.0%; Pred
0; N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY32836 standard; protein; 282
                                                                                                                                                                                                                                                                                                                    seqdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-00021769.
98US-0074294P.
98US-0105509P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Woodchuck hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                           6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Punnonen J, Bass SH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infections and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-518452/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MAXY-) MAXYGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   265 WRWLRR 270
                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 WRWLRR 15
                                                                                                                                                                                                                                                                                                                                                              Sequence 277 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB; AAZ10971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09941383-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY32836
AAY32836
AAY32836
AAY3
AAY
AAY32836
AAY3
AAY32836
AAY3
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AAY33836
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AAY3386
AAY36

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                                                                                                                                                                                                                                                                                                                                                                       The invention comprises the amino acid sequence of proteins from the hyperthermophile Methanopyrus kandleri, the invention also comprises the complete genome from Methanopyrus kandleri. The Methanopyrus kandleri proteins of the invention are useful for enhancing the stability and/or activity of other proteins. The Methanopyrus kandleri genome is useful in a variety of diagnostic and analytical methods. The present amino acid sequence represents a Methanopyrus kandleri protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and
                                                                                                                                                                                                                                   New isolated nucleic acids encoding any of about 1700 Methanopyrus kandleri proteins, and the encoded proteins, useful as a medicaments or as diagnostic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment c pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 7; Length 277; 5. 1.7e+02;
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                                                                                                                       Kozyavkin S;
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                                                                                                                                                                                                                                                                                                                                   Claim 31; SEQ ID NO 1353; 1023pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 6; I
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pseudomonas aeruginosa polypeptide #996.
                                                                                                                       Pavlova N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30.0%;
  16-SEP-2002; 2002US-0410974P.
                                                FIDELITY SYSTEMS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Conservative
                                                                                                                     Slesarev AI, Pavlov A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pseudomonas aeruginosa
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 WRWLRR 15
                                                                    MALYKH A
                                                                                                                                                                                       N-PSDB; ADM27081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 277 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; ABD02392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rubenfield MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6551795-B1
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2000US-0225270P.

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14-SEP-2000; 2000US-0232399P.
14-SEP-2000; 2000US-0232400P.
14-SEP-2000; 2000US-0232400P.
14-SEP-2000; 2000US-0233064P.
14-SEP-2000; 2000US-0233064P.
11-SEP-2000; 2000US-0233064P.
21-SEP-2000; 2000US-023423P.
21-SEP-2000; 2000US-023499P.
22-SEP-2000; 2000US-023499P.
22-SEP-2000; 2000US-0236494P.
27-SEP-2000; 2000US-0236494P.
27-SEP-2000; 2000US-0236349P.
29-SEP-2000; 2000US-023634P.
29-SEP-2000; 2000US-023636P.
29-SEP-2000; 2000US-023636P.
29-SEP-2000; 2000US-023636P.
29-SEP-2000; 2000US-023636P.
29-SEP-2000; 2000US-023636P.
29-SEP-2000; 2000US-023636P.
29-CCT-2000; 2000US-023636P.
29-CCT-2000; 2000US-023636P.
29-CCT-2000; 2000US-023638P.
29-CCT-2000; 2000US-023638P.
20-CCT-2000; 2000US-023636P.
20-CCT-2000; 2000US-0237039P.
20-CCT-2000; 2000US-0237039P.
20-CCT-2000; 2000US-0237039P.
20-CCT-2000; 2000US-0237039P.
20-CCT-2000; 2000US-0237039P.
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01-SEP-2000; 2000US-0229344P.

05-SEP-2000; 2000US-0229509P.

05-SEP-2000; 2000US-0229513P.

05-SEP-2000; 2000US-0239513P.

06-SEP-2000; 2000US-023043P.

08-SEP-2000; 2000US-0231242P.

08-SEP-2000; 2000US-0231244P.

08-SEP-2000; 2000US-0231244P.

08-SEP-2000; 2000US-0231244P.

08-SEP-2000; 2000US-0231244P.

08-SEP-2000; 2000US-0231244P.
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22-AUG-2000; 2000US-02266BLP.
22-AUG-2000; 2000US-022668PP.
22-AUG-2000; 2000US-0227182P.
23-AUG-2000; 2000US-0227009P.
                                                                                                                                                                     14-AUG-2000; 2000US-0225447P.
14-AUG-2000; 2000US-0225757P.
14-AUG-2000; 2000US-0225758P.
14-AUG-2000; 2000US-0225759P.
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20-OCT-2000;
20-OCT-2000;
20-OCT-2000;
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08-NOV-2000;
08-NOV-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
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01-NOV-2000;
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related antigenic polypeptide. The library can be used to produce a recombinant multivalent antigenic polypeptides of the invention, that contains at least two antigenic determinants (AD) from different polypeptides. The multivalent antigenic polypeptides are used in vaccines to induce a protective or therapeutic response to a wide variety of infectious agents (bacteria, viruses, parasites, including Plasmodium falciparum); allergens, asthma; autoimmune disease (e.g. rheunatoid ard canner, also, where directed against sperm antigens, they can be used for contraception. The multivalent peptides can be evolved to induce an optimised immune response against a wide variety of antigens, particularly a broad spectrum response to many different strains of a pathogen, including strains that are likely to appear in the future. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Central nervous system; CNS; autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; ischaemia; angiogenesis; nervous system disorder; Alzheimer's disease; AIDS; ocular disorder; acquired immunodeficiency virus; dysphagia; gastrointestinal disorder; adenocarcinoma; reproductive system disorder; testicular feminisation; endocrine disorder; diabetes; cancer; leukaemia; neovascularisation; respiratory disorder; renal disorder; kidney failure; blood disorder; myocardial infarction; wound healing; cell proliferation; skin aging; food additive; food preservative; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                    30.0%; Score 6; DB 2; Length 282; 100.0%; Pred. No. 1.7e+02; ative 0; Mismatches 0; Indels
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2000US-0205515P
2000US-0209467P
2000US-0215135P
2000US-0215135P
2000US-021680P
2000US-021680P
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2000US-0180628P.
2000US-0184664P.
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2000US-0189874P.
2000US-0190076P.
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2000US-0220963P.
2000US-0220964P.
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                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 100.
es 6, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      210 SSWALG 215
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                                                                                                                                                                                                                                                                                                                                Sequence 282 AA;
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16-MAR-2000;
17-MAR-2000;
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28-JUN-2000;
30-JUN-2000;
07-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-FEB-2000;
24-FEB-2000;
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Matches
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2000US-0231414P. 2000US-0232080P. 2000US-0232081P. 2000US-0232397P. 2000US-0232397P.

2000US-0246475P 2000US-0246476P

2000US-0241808P

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The invention describes an isolated nucleic acid molecule (I) encoding a novel central nervous system protein. (I) and polypeptides (III) encoded by (I), are used to treat a medical conditions and in diagnosis of a pathological condition. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. carebral ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease and amylotrophic lateral sclerosis, infections caused by bacteria, viruses e.g. Acquired immunodeficiancy virus (AIDS) and fungi, ocular disorders e.g. Acquired immunodeficiancy virus (AIDS) and fungi, ocular disorders e.g. corneal infection, gastrointestinal disorders e.g. dysphagia, adenocarcinomas and irritable bowel syndrome, reproductive system disorders e.g. testicular feminisation, endocrine disorders e.g. diabetes and pituitary dwarfism, cancers and disorders at the cellular level e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid encoding a protein for diagnosing, preventing, treating or ameliorating medical conditions and used as food additives or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 9; SEQ ID NO 621; 837pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-0249300P.
200US-0250160P.
200US-0250391P.
200US-0251030P.
2000US-025198P.
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2000US - 0246613P
2000US - 0249207P
2000US - 0249208P
2000US - 0249210P
2000US - 0249211P
2000US - 0249216P
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2000US-0249265P.
2000US-0249297P.
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2000US-0251856P.
                                                                                                                    2000US-0246532P.
2000US-0246609P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      000US-0249299P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0251868P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   preservatives.
                                                                                    08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
                                                                                                                                                08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
                                                        08-NOV-2000;
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17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                              17-NOV-2000;
17-NOV-2000;
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17-NOV-2000;
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Authorical disease; Managorian, garacological; gastrointestinal; respiratory; antiparasitic; muscular; gynaecological; gastrointestinal; respiratory; ardivascular; antiarteriosclerotic; antiarrhythmic; cardiant; nephrotropic; litholytic; cytostatic; gene therapy; neural disorder; Alzheimer's disease; Parkinson's disease; Huntington's chorea; wamyotrophic lateral sclerosis; multiple sclerosis; multiple sclerosis; myotrophic lateral sclerosis; multiple sclerosis; inflammatory disorder; dabetes; rhematorid arthritis; systemic lupus erythematosus; autoimmune thyroiditis; haemolytic anaemia; systemic lupus erythematosus; crohn's disease; inflectious disease; inflammatory disorder; ischaemia-reperfusion injury; inflammatory bowal disease; Crohn's disease; inflection; bacterial infection; fungal infection; parasitic infection; bacterial infection; fungal infection; parasitic infection; muscular disorder; gastrointestinal disorder; pulmonary disorder; renal disorder; acute glomerulonephritis; pyelonephritis; renal disorder; acute glomerulonephritis; pyelonephritis; renal lithiasis; proliferative disorder; cancerous diseases; human.
                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antidiabetic, antirheumatic, antiarthritic, dermatological,
antiinflammatory, immunosuppressive, antithyroid, antianemic, vasotropic,
anti-HIV, hepatotropic, virucide, antibacterial; fungicide,
leukaemia, disorders involving neovascularisation e.g. malignancies, respiratory disorders e.g. nonallergic rhinitis, renal disorders e.g. acute kidney failure and blood related disorders e.g. myocardial infarction. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, protein,
                                                                                                                                                                                                                                                              .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroprotective; nootropic; antiparkinsonian; anticonvulsant;
                                                                                                                                                                                                                  Length 305;
                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                30.0%; Score 6; DB 4; Ler 100.0%; Pred. No. 1.8e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADI54418 standard; protein; 305 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human protein seq id 621.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-MAR-2000; 2000US-0189874P.
17-MAR-2000; 2000US-0190076P.
18-APR-2000; 2000US-0198123P.
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2000US-0180628P.
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2000US-0186350P.
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2000US-0209467P.
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2000US-0215135P.
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2000US-0216880P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0217487P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-APR-2004 (first entry)
                                                                                                                                                                                                                                                            6; Conservative
                                                                                                                                                                                                                                                                                                                                             216 WLRRYG 221
                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                   12 WLRRYG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-MAY-2000;
07-JUN-2000;
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ADI54418
                                                                                                                                                                                                                                                            Matches
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2000US-0218290P

14-JUL-2000;

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PR 26-JUL-2000; 2000US-02220564P.
PR 14-ANG-2000; 2000US-02225114P.
PR 14-ANG-2000; 2000US-0225514P.
PR 14-ANG-2000; 2000US-0225514P.
PR 14-ANG-2000; 2000US-0225514P.
PR 14-ANG-2000; 2000US-022554P.
PR 14-ANG-2000; 2000US-022556P.
PR 14-ANG-2000; 2000US-022556P.
PR 14-ANG-2000; 2000US-022556P.
PR 14-ANG-2000; 2000US-022556P.
PR 14-ANG-2000; 2000US-022557P.
PR 14-ANG-2000; 2000US-022556P.
PR 14-ANG-2000; 2000US-022557P.
PR 14-ANG-2000; 2000US-022575P.
PR 14-ANG-2000; 2000US-022579P.
PR 14-ANG-2000; 2000US-022575P.
PR 22-ANG-2000; 2000US-022579P.
PR 22-ANG-2000; 2000US-022944P.
PR 15-SEP-2000; 2000US-022944P.
PR 15-SEP-2000; 2000US-022944P.
PR 05-SEP-2000; 2000US-022944P.
PR 05-SEP-2000; 2000US-022944P.
PR 06-SEP-2000; 2000US-022944P.
PR 06-SEP-2000; 2000US-022944P.
PR 06-SEP-2000; 2000US-022944P.
PR 14-SEP-2000; 2000US-022944P.
PR 14-SEP-2000; 2000US-022944P.
PR 14-SEP-2000; 2000US-023949P.
PR 15-SEP-2000; 2000US-023949P.
PR 25-SEP-2000; 2000US-
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R 08-NOV-2000; 2000US-0246478R 08-NOV-2000; 2000US-0246478R 08-NOV-2000; 2000US-0246478R 08-NOV-2000; 2000US-0246779R 08-NOV-2000; 2000US-0246729R 08-NOV-2000; 2000US-0246229R 08-NOV-2000; 2000US-0249208R 17-NOV-2000; 2000US-0249208-

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GPCR; G protein coupled receptor; signal transduction; olfactory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 10; SEQ ID NO 4; 97pp + Sequence Listing; Japanese.
                                                                                                                                                                         drug development; gustatory; taste; fragrance; receptor
                     ABP95597 standard; protein; 368 AA.
                                                                                                                        Human GPCR polypeptide SEQ ID NO 4.
                                                                                                                                                                                                                                                                                                                                                                                             (NISC-) JAPAN SCI & TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                              Mitaku S;
                                                                                                                                                                                                                                                                                                                                                           13-FEB-2001; 2001JP-00034434.
                                                                                                                                                                                                                                                                                                          30-JUL-2001; 2001WO-IB001446
                                                                                                                                                                                                                                                                                                                                           04-AUG-2000; 2000JP-00237818
                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-304118/34.
N-PSDB; ABZ42871.
                                                                                                                                                                                                                                                                                                                                                                                                                              Takeda S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                         WO200216548-A2
                                                                                                                                                                                                           Homo sapiens.
                                                                                      06-MAR-2003
                                                                                                                                                                                                                                                                          28-FEB-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                             Haga T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 31
     ABP95597
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Huntington's chorea, amyotrophic lateral sclerosis or multiple sclerosis, immune system disorders, e.g. diabetes, rheumatoid arthritis, systemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasmid pCN4 carries an insert (AAT49596) coding for a fusion (AAW09046) between the woodchuck hepatitis virus (WHV) core protein amino acids 1-177 with amino acid 47 of the WHV small surface protein. Corransfection of human hepatoma HuH-7 cells with wild-type WHV pCWM82 and mutant core construct pCN4 resulted in an 85% inhibition of wild-type viral DNA toxic to HCC cells. The dominant negative core polypeptide was not toxic to HCC cells. The dominant negative effects on viral replication are consistent over a range of hepathavirus species and can be utilised in the treatment of viral infection. (Updated on 17-OCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                       Hepadnavirus, hepatitis B virus, HBV; hepatitis delta virus,
ground squirrel hepatitis B virus, duck hepatitis B virus, core protein;
                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 introducing a mutant polypeptide with a mutated core protein or corresponding nucleic acid, for treating, e.g. hepatitis B.
                                                                                      0;
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0
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                                      DB 8; Lt.,
40. 1.8e+02;
0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 2; Length 346;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                        WHV core-surface fusion dominant negative polypeptide.
                                               30.0%; Score 6; DB 8
100.0%; Pred. No. 1.8
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30.0%; Score 6; DB 2
100.0%; Pred. No. 2e+
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           replication; antiviral; therapy; pCN4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Melegari M;
                                                                                                                                                                                                                                     AAW09046 standard; protein; 346 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96WO-US010602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95US-0017814P
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                                                                                                                                                                                                                                                                                                                        (first entry)
                                 Query Match
Best Local Similarity 100...
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                                                                                                                                                                                                                                                                                                        (revised)
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                                                                                                                                                  216 WLRRYG 221
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Best Local Similarity
Matches 6; Conserv
                                                                                                                    12 WLRRYG 17
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11-APR-1997
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ID AAW0
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CORP.

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The invention relates to a method for screening G protein-coupled receptor (GPCR) genes (ABZ42870-ABZ43216) and/or GPCR proteins (ABP95596-ABP95421) by extracting open-reading frames containing 6-8 transmembrane domains with 250-1000 amino acid residues to give a gene homologous with a known GPCR gene. The receptor proteins and encoded genes are useful for studying in vivo signal transduction mechanism and identifying targets for drug development e.g. based on olfactory and gustatory receptors in form of agonists and antagonists by screening intrinsic and extrinsic ligands as bitter taste inhibitors, taste enhancers and fragrance improvers. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence of gpG encoded by segment of Xanthomonas campestris DNA that contains a gene cluster that directs Xanthan biosynthesis.
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to, 2.1e+02;
0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; DB 5
100.0%; Pred. No. 2.1
iive 0; Mismatches
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13-FEB-1991
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RESULT 30

SSWALG 9

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Disclosure; SEQ ID NO 7940; 932pp; English.
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Carr GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAR-2001; 2001US-00815242.
06-SEP-2001; 2001US-0094893.
25-OCT-2001; 2001US-0342923P.
08-FEB-2002; 2002US-00372851.
06-MAR-2002; 2002US-0362699P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2002; 2002WO-US009107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUN-2003 (first entry)
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                                                                                                                                                                                                                                                                                           6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zamudio C,
Trawick JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-029926/02
                                                                                                                                                                                                                                                                                                                                                                       388 WLRRYG 393
                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                              12 WLRRYG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Borrelia cepacia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; ACA23802
                                                                                                                                                                                                                Sequence 395 AA;
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Wall D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABU19932;
                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid encoding a Klebsiella pneumoniae polypeptide, useful for preparing a vaccine composition against Klebsiella pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         contains
                                                                                                                                                                                                                                                                                                                                                   variants - by transforming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Virtually all of the segment of Xanthomonas campestris DNA that contain a gene cluster that directs Xanthan biosynthesis (AAN70753), codes for protein products. Each gene is designated by a letter (see Fig 11) and its protein product is designated by that letter preceded by '9p' (AAP70455-67). (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                   Recombinant DNA prodn. of xanthan gum or its variants - by trans! host cells with vector contg. DNA coding for enzymes involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recombinant expression vector; transcription regulatory element; Klebsiella pneumoniae protein; antibacterial; Vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.0%; Score 6; DB 1; Length 379; 100.0%; Pred. No. 2.2e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                         Vanderslic RW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Klebsiella pneumoniae polypeptide segid 7940,
                                                                                                                                                                                                                                                                         Betlach MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO61423 standard; protein; 395 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENO-) GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                              Example, Fig 12; 149pp; English.
                                                                                                                                  87WO-US000604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JAN-2000; 2000US-00489039
                                                                                                                                                                         86US-00842944
                                                                                                                                                                                           87US-00029530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0117747P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                         polysaccharide synthesis.
                                                                                                                                                                                                                                  (GETT-) GETTY SCI DEV CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Conservative
                                                                                                                                                                                                                                                                         Doherty DH,
                  Xanthomonas campestris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Osborne M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Klebsiella pneumoniae
                                                                                                                                                                                                                                                                                                            WPI; 1987-291651/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-895346/82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            279 GWRWLR 284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9 GWRWLR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; ACH94974.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 379 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JAN-1999;
                                                                                                                                  24-MAR-1987;
                                                                                                                                                                       24-MAR-1986;
23-MAR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US6610836-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JUL-2004
                                                                                             08-OCT-1987
                                                                                                                                                                                                                                                                         Capage MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Breton GL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 32
ABO61423
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Matches
PAY X PAY X PAY Y PAY X 
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(1) a vector comprising a promoter operably linked to the nucleic acid encoding a polypeptide whose expression is inhibited by the antisense morteic acid; (2) a host cell containing the vector; (3) an isolated polypeptide or its fragment whose expression is inhibited by the antisense antisense nucleic acid; (4) an antibody capable of specifically binding the polypeptide; (6) producing the polypeptide; (6) inhibiting callular proliferation or the activity of a gene in an operon required for proliferation; (7) identifying a compound that influences the activity of
The invention describes a new isolated nucleic acid encoding a Klebsiella premoniae polypeptide. Also described are: a recombinant expression vector comprising the nucleic acid, operably linked to a transcription regulatory element; and a cell comprising the recombinant expression vector. The nucleic acid is useful for preparing a vaccine composition against Klebsiella pneumoniae. This is the amino acid sequence of a Klebsiella pneumoniae, and the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense nucleic acids, useful for identifying proteins or screening for homologous nucleic acids required for cellular proliferation to isolate candidate molecules for rational drug discovery programs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense; prokaryotic essential gene; cell proliferation; drug design.
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Xu HH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .;
o
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 395;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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Forsyth RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein encoded by Prokaryotic essential gene #5459.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 7; Ler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Haselbeck R,
Yamamoto R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 25; SEQ ID NO 47856; 1766pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  30.0%; Score 6; DB 7
100.0%; Pred. No. 2.2
iive 0; Mismatches
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03-JUN-2003; 2003US-00454246.

(CURA-) CURAGEN CORP.

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0
                                                                                                                                                                                                                drug discovery programs, or for screening homologous nucleic acids required for proliferation in cells other than S. aureus, S. typhimurium, K. pneumoniae or P. aeruginosa. The present sequence is encoded by one of
                                                                                               compound's activity; (11) a cullure comprising stains in which the gene product is overexpressed or underexpressed; (12) determining the extent to which each of the strains is present in a culture or collection of strains; or (13) identifying the target of a compound that inhibits the strains; or (a) identifying the target of a compound that inhibits the identifying proteins. The antisense nucleic acids are useful for identifying proteins or screening for homologous nucleic acids required for cellular proliferation to isolate candidate molecules for rational
                                                                                                                                                                                                                                                               the target prokaryotic essential genes. Note: The sequence data for this spatent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published_pct_sequences
the gene product or that has an activity against a biological pathway required for proliferation, or that inhibits cellular proliferation; (8) identifying a gene required for cellular proliferation or the biological pathway in which a proliferation-required gene or its gene product lies or a gene on which the test compound that inhibits proliferation of an organism acts; (9) manufacturing an antibiotic; (10) profiling a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; NOVX; secreted protein; cancer; diabetes; obesity;
endocrine disorder; CNS disorder; inflammatory disorder; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                   Length 396;
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                                                                                                                                                                                                                                                                                                                                                                                 DB 6; Len
                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; DB 6
100.0%; Pred. No. 2.3
Live 0; Mismatches
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2002US-0386641P.
2002US-0386643P.
2002US-0386416P.
2002US-0386816P.
2002US-0387002P.
2002US-0387659P.
2002US-0387659P.
2002US-0387534P.
2002US-0389723P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002US-0396706P
2002US-0402832P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002US-0403522P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002US-0387037P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 ALGWRW 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               75 ALGWRW 80
                                                                                                                                                                                                                                                                                                                                                  Sequence 396 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004000997-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-JUN-2002;
13-JUN-2002;
17-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JUL-2002;
12-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-AUG-2002;
06-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ34131;
                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 34
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protein comprising one or more conservatives, their mature lorms, a protein comprising on or more conservatives, their mature lorms, a least 95% identity one of the 141 proteins. Also included are a composition comprising NOVX (or a NOVX nucleic acid molecule (NA)), a kit comprising the composition of NOVX incleic acid molecule encoding a NOVX protein, producing NOVX (comprising culturing a cell under conditions that lead to expression of the culturing a cell under conditions that lead to expression of the coll comprises a vector comprising NOVX (comprising CC polypeptide, where the call comprises a vector comprising NOVX NA), identifying an agent that binds to NOVX, identifying a potential condition or physiological interactions of NOVX, screening for a modulator of activity of or latency or predisposition to a comprising a pathology associated with NOVX, treating a pathological confined activity of NOVX, treating or preventing a pathology associated with NOVX modulating the activity of NOVX, treating call comprising the vector comprising the NOVX moduleic acid molecule, a coll comprising the presence or amount of NOVX or the nucleic acid molecule in a sample, and determining the presence of properior of properior or predisposition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ·.
                                                                                                                                                                         Stone DJ;
Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        compositions of the present invention are useful for the diagnosis and treatment of disorders associated with aberrant expression or activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     molecule in a sample, and determining the presence of or predisposition to a disease associated with altered levels of expression of NOVX or the nucleic acid molecule in a first mammalian subject. The methods and
                                                                                  SJ, Edinger SR;
Guo X, Gusev VY, Ji W;
, Ort T, Padigaru M;
                                                                                                                                                                                                                                                                                                     New isolated NOVX polypeptides useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory or CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the NOVX polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing, gene therapy and predictive medicine. The present sequence represents a NOVX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis B virus, HBV, preCore, Core, preS1, preS2, HBS, HBX, HBPol, HBsAg, antiviral, vaccine, genotype G, genotyping, HBcAg, HBeAg.
                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a new isolated polypeptide (designated NOVX) comprising one of 141 fully defined sequences, their mature forms, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                       Gorman L, Guo X, Gusev VY
, Millet I, Ort T, Padiga
A, Peyman JA, Rieger DK;
G, Smithson G, Spytek KA,
AM, Voss EZ, Zerhusen BD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30.0%; Score 6; DB 8; Len 100.0%; Pred. No. 2.3e+02; iive 0; Mismatches 0;
                                                                                       Casman SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HBV genotype F preS1/preS2/HBsAg polypeptide.
                                                                                                       Gerlach VL, Gorm
Malyankar UM, Mi
an M, Pena CEA, F
ee P, Shenoy SG, S
                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 20; 418pp; English.
                                                                                       Burgess CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAG66933 standard; protein; 400 AA.
                                                                                                                             Li L, Macdougall JR, Maly
Prayaga SK, Patturajan M,
Rothenberg ME, Sciore P,
Taupier RJ, Tchernev VT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-2001 (first entry)
                                                                                       W, Boldog FL,
Ellerman K, Ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; Conservative
                                                                                                                                                                                                                                           WPI; 2004-082483/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58 WSSWAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 WSSWAL
                                                                                       Anderson DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAG66933;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                Eisen A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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ΩD
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Homo sapiens
        05-JUL-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
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                                                                                                                                                                                                                                                           The invention relates to the complete nucleic acid sequence of a new human hepatitis B virus (HBV) genotype, provisionally named genotype G. This genotype was found with a high prevalence in patients chronically infected with HBV and residing in Burope and the USA. The invention relates to a fully defined sequence of 3248 mucleotides as given in sequence that is degenerate to the mentioned sequences. These polymucleotides are useful for HBV genotyping. The proteins encoded by the polymucleotides are useful for detecting antibodies in a biological sample. Ligands that bind to the proteins and antibodies directed against the proteins are useful for detecting the proteins and for detecting the proteins and surfaced against the proteins are useful for detecting the proteins and solventing and preparing a vaccine or medicament for treating HBV infections. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                        present sequence is provided in an amino acid sequence alignment of the preS1, preS2 and HBsAg open reading frame of the different HBV genotypes
                                                                                                                                                                                              HBV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           molecule protein, MOL protein, MOLX, MOLX agonist, MOLX antagonist, cardiant, antidiabetic, antiarteriosclerotic, gene therapy, MOLX-associated disorder, cardiomyopathy, diabetes, atherosclerosis,
                                                                                                                                                                                  Novel isolated and/or purified hepatitis B virus polypeptide and polynucleotide sequences that are phylogenetically different from genotype A-F molecules, useful for HBV diagnosis, prophylaxis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 400;
5. 2.3e+02;
cches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; DB 4
100.0%; Pred. No. 2.3
:ive 0; Mismatches
                                                                                                                                         De Gendt S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD18225 standard; protein; 401 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human molecule (MOL) protein MOL4d.
                                                                                                                                                                                                                                          Example 3; Fig 6; 94pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JUL-2002; 2002WO-US021268.
                                                               20-NOV-2000; 2000WO-EP011526
                                                                                   99EP-00870252
99US-0169287P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match
Best Local Similarity 100...
6; Conservative
                                                                                                                                         Van Geyt C,
                                                                                                                    (INNO-) INNOGENETICS NV
                                                                                                                                                              WPI; 2001-374785/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                328 SSWALG 333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           σ
Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 400 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 SSWALG
                    WO200140279-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; MOL4d
                                                                                     03-DEC-1999;
                                                                                               07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JAN-2003
                                                                                                                                           Stuyver L,
                                          07-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD18225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 36
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novel molecule (MOL) proteins numbered MOLI-23, referred to generally in the specification as MOLX. Compounds which modulate the function of the MOLX proteins of the invention, MOLX agonists or antagonists, may have cardiant, antidiabetic or antiatrateriosclerotic activities. In addition, the DNA and protein sequences disclosed may prove useful for gene therapy. The protein, nucleic acid or antibody is useful for gene composition for treating or preventing a MOLX-associated disorder, for example cardiomyopathy, diabetes or atherosclerosis. The present sequence is the amino acid sequence of a MOL protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , Vernet CAM, Shimkets RA, Anderson DW, Padigaru M;
ii, Shenoy SG, Casman SJ, Rastelli L, Alsobrook JP;
Grosse WM, Gusev VY, Ji W, Lepley DM, Liu X, Mezick AJ;
Shen L, Spaderna SK, Spytek KA, Szekeres ES;
Tchernev VT, Zerhusen BD, Voss EZ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New MOLX polypeptide, nucleic acid or MOLX-specific antibody, useful preparing a composition for treating or preventing a MOLX-associated disorder, e.g., cardiomyopathy, diabetes or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   molecule protein; MOL protein; MOLX; MOLX agonist; MOLX antagonist;
cardiant; antidiabetic; antiarteriosclerotic; gene therapy;
MOLX-associated disorder; cardiomyopathy; diabetes; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to novel human nucleic acid sequences which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 7; Length 401;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.0%; Score 6; DB 7
100.0%; Pred. No. 2.3
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 111; 371pp; English.
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                        05-JUL-2001; 2001US-0303241P.
26-SEP-2001; 2001US-0303241P.
26-SEP-2001; 2001US-0096545.
26-SEP-2001; 2001US-00966546.
01-APR-2002; 2002US-0368996P.
01-APR-2002; 2002US-0368996P.
08-MAY-2002; 2002US-0368936P.
07-JUN-2002; 2002US-0384327P.
07-JUN-2002; 2002US-0386816P.
2001US-0303168P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Boldog FL, Li L, W., Gur
Burgess CE, Grosse WM, Gur
Burgess CE, Grosse WM, Gur
Patturajan M, Shen L, Spa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-210304/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fernandes ER, Vern
Boldog FL, Li L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 WSSWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 401 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; ADD18223
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05-JUL-2001; 2001US-0303241P.
26-SEP-2001; 2001US-00965212.
26-SEP-2001; 2001US-00966545.
26-SEP-2002; 2001US-0966546.
01-APR-2002; 2002US-0368996P.
01-APR-2002; 2002US-0369965P.
08-MAY-2002; 2002US-038937379.
06-MAY-2002; 2002US-0384327P.
07-JUN-2002; 2002US-0384327P.
                        03-JUL-2002; 2002WO-US021268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 WSSWAL 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 WSSWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 401 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003003984-AZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; MOL4g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 39
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    à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       novel molecule (MOL) proteins numbered MOLI-23, referred to generally in the specification as MOLX. Compounds which modulate the function of the MOLX proteins of the invention, MOLX agonists or antagonists, may have cardiant, antidiabetic or antazeric/sclerotic activities. In addition, the DNA and protein sequences disclosed may prove useful for gene composition for treating or preventing a MOLX-associated disorder, for example cardiomyopathy, diabetes or atherosclerosis. The present sequence is the amino acid sequence of a MOL protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                               Mezick AJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New MOLX polypeptide, nucleic acid or MOLX-specific antibody, useful for preparing a composition for treating or preventing a MOLX-associated disorder, e.g., cardiomyopathy, diabetes or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                       Padigaru M;
Alsobrook JP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             molecule protein; MOL protein; MOLX; MOLX agonist; MOLX antidiabetic; antiarteriosclerotic; gene therapy;
MOLX-associated disorder; cardiomyopathy; diabetes; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to novel human nucleic acid sequences which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
                                                                                                                                                                                                                                                                                                                                   ER, Vernet CAM, Shimkets RA, Anderson DW, Padiga:
, Li L, Shency SG, Casman SJ, Rastelli L, Alsobrv
E, Grosse WM, Gusev VY, Ji W, Lepley DM, Liu X,
M, Shen L, Spaderna SK, Spytek KA, Szekeres ES;
J, Tchernev VT, Zerhusen BD, Voss EZ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30.0%; Score 6; DB 7; Length 401; 100.0%; Pred. No. 2.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 116; 371pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human molecule (MOL) protein MOL4e.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD18228 standard; protein; 401 AA
                                     2001US-0303168P.
2001US-0303241P.
2001US-00965212.
2001US-00966545.
2001US-00966546.
                                                                                                                                           2002US-0368996P.
2002US-0369065P.
2002US-0378730P.
                                                                                                                                                                                                         2002US-0384327P
2002US-0386816P
  03-JUL-2002; 2002WO-US021268
                                                                                                                                                                                                                                                    2002US-00174372
                                                                                                                                                                                                                                                                                                                                                       Boldog FL, Li L, Shenoy,
Burgess CE, Grosse WM, G
Patturajan M, Shen L, Sp
Taupier RJ, Tchernev VT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Conservative
                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2003-210304/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 WSSWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-210304/
N-PSDB; ADD18229.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 401 AA;
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                                     05-JUL-2001; 2
05-JUL-2001; 2
26-SEP-2001; 2
26-SEP-2001; 2
26-SEP-2001; 2
                                                                                                                                           01-APR-2002; 2
01-APR-2002; 2
08-MAY-2002; 2
                                                                                                                                                                                                           30-MAY-2002;
07-JUN-2002;
                                                                                                                                                                                                                                                    17-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                       Fernandes
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novel molecule (WOL) proteins numbered MOL1-23, referred to generally in the specification as MOLX. Compounds which modulate the function of the MOLX proteins of the invention, MOLX agonists or antagonists, may have cardiant, antidiabetic or antidazeric or attidiate or attidiate or attidiate or attidiate or attidiate or attidiate or attidiate. In addition, the DNA and protein sequences disclosed may prove useful for gene therapy. The protein, nucleic acid or antibody is useful for preparing a composition for treating or preventing a MOLX-associated disorder, for example cardiomyopathy, diabetes or atherosclerosis. The present sequence is the amino acid sequence of a MOL protein of the invention.
                                                                                                                                                               Fernandes ER, Vernet CAM, Shimkets RA, Anderson DW, Padigaru M; Boldog FL, Li L, Shenoy SG, Casman SJ, Rastelli L, Alsobrook JP; Burgess CB, Grosse WM, Gusev VY, Ji W, Lepley DM, Liu X, Mezick AJ; Patturajan M, Shen L, Spaderna SK, Spytek KA, Szekeres ES; Taupier RJ, Tchernev VT, Zerhusen BD, Voss EZ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to novel human nucleic acid sequences which encode
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New MOLX polypeptide, nucleic acid or MOLX-specific antibody, useful for preparing a composition for treating or preventing a MOLX-associated disorder, e.g., cardiomyopathy, diabetes or atherosclerosis.
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cardiant; antidiabetic; antiarteriosclerotic; gene therapy;
MOLX-associated disorder; cardiomyopathy; diabetes; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.0%; Score 6; DB 7
100.0%; Pred. No. 2.3
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 114; 371pp; English.
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17-JUN-2002; 2002US-00174372
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                                                                                         (CURA-) CURAGEN CORP.
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N-PSDB; ADD18226.
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Fernandes ER, Vernet CAM, Shimkets RA, Anderson DW, Padigaru M, Boldog FL, Li L, Shenoy SG, Casman SJ, Rastelli L, Alsobrook JP; Burgess CE, Grosse WM, Gusev VY, Ji W, Lepley DM, Liu X, Mezick AJ; Patturajan M, Shen L, Spadera SK, Spytek KA, Szekeres ES; Taupier RJ, Tchernev VT, Zerhusen BD, Voss EZ,
                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to novel human nucleic acid sequences which encode
                                                                                                                                                                                                                                                                                                                                          New MOLX polypeptide, nucleic acid or MOLX-specific antibody, useful for preparing a composition for treating or preventing a MOLX-associated disorder, e.g., cardiomyopathy, diabetes or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 118; 371pp; English.
                                                        200105-0303241F.
200105-0365545.
200105-0966546.
200105-0966546.
200205-0368966F.
200205-036996F.
200205-034937P.
200205-0349437P.
200205-0346916P.
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Boldog FL, ...
Burgess CB, Grosse ...
--turajan M, Shen L, Spau...
--turajan T Tchernev VT, Z
                      03-JUL-2002; 2002WO-US021268
                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                      WPI; 2003-210304/20.
                                                                                                                                                                                                                                                                                                                   N-PSDB; ADD18231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 401 AA;
                                                          05-JUL-2001;
26-SEP-2001;
26-SEP-2001;
26-SEP-2001;
01-APR-2002;
01-APR-2002;
                                                                                                                                    08-MAY-2002;
30-MAY-2002;
                                                                                                                                                             07-JUN-2002;
17-JUN-2002;
                                               05-JUL-2001;
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novel molecule (MOL) proteins numbered MOL1-23, referred to generally in the specification as MOLX. Compounds which modulate the function of the MOLX proteins of the invention, MOLX agonists or antagonists, may have cardiant, antidiabetic or antiatreticolerchic activities. In addition, the DNA and protein sequences disclosed may prove useful for gene composition for treating or preventing a MOLX-associated disorder, for example cardiomyopathy, diabetes or atherosclerosis. The present sequence is the amino acid sequence of a MOL protein of the invention.
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Matches
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ADJ34125 standard; protein; 401 AA.
                                 60 WSSWAL 65
3 WSSWAL 8
                                                                                      RESULT 40
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Human, NOVX; secreted protein; cancer; diabetes; obesity; endocrine disorder; CNS disorder; inflammatory disorder; gene therapy.
                                                     Human secreted protein NOV2e
                                    06-MAY-2004 (first entry)
                                                                                                     Homo sapiens
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WO2004000997-A2

31-DEC-2003

Sequence 401 AA;

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Stone DJ;
Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                      New isolated NOVX polypeptides useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory or CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the NOVX polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing, gene
                                                                                                                                                                                                                                                                                                       Guo X, Gusev VY, Ji
, Ort T, Padigaru M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy and predictive medicine. The present sequence represents a
                                                                                                                                                                                                                                                                                          Anderson DW, Boldog FL, Burgess CE, Casman SJ, Edinger SR, Eisen A, Ellerman K, Gerlach VL, Gorman L, Guo X, Gusev VY Li L, Macdougall JR, Malyankar UM, Millet I, Ort T, Padiga Prayaga SK, Patturajan M, Pena CEA, Peyman JA, Rieger DK, Rothenberg ME, Sciore P, Shenoy SG, Smithson G, Spytek KA, Taupier RJ, Tchernev VT, Vernet CAM, Voss EZ, Zerhusen BD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 14; 418pp; English.
                                                                               07-JUN-2002; 2002US-0386816P.

07-JUN-2002; 2002US-0387002P.

10-JUN-2002; 2002US-0387640P.

11-JUN-2002; 2002US-0387659P.

12-JUN-2002; 2002US-0387934P.

13-JUN-2002; 2002US-0389723P.

17-JUN-2002; 2002US-0389723P.
                                                  2002US-0386041P.
2002US-0386453P.
                                                                                                                                                                                                      13-AUG-2002; 2002US-0403486P.
14-AUG-2002; 2002US-0403522P.
15-AUG-2002; 2002US-0403748P.
06-NOV-2002; 2002US-0403748P.
       04-JUN-2003; 2003WO-US017512
                                                                                                                                                             2002US-0389742P
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12-AUG-2002; 2002US-0402832P
                                                                        2002US-0386974P
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                                                                       06-JUN-2002;
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06-JUN-2002;
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protein comprising one or more conservative substitutions or having at least 95% identity to one of the 141 proteins. Also included are a composition comprising NOVX (or a NOVX nucleic acid molecule (NA)), a kit comprising the composition of NOVX in one or more containers, an isolated nucleic acid molecule encoding a NOVX protein, producing NOVX (comprising culturing a cell under conditions that lead to expression of the polypeptide, where the cell comprises a vector comprising NOVX MA), identifying an agent that binds to NOVX, identifying a potential comprises a vector comprising NOVX MA, identifying a potential conspiculation or a mediator of activity of or latency or predisposition to a pathology associated with NOVX, modulating the activity of NOVX, treating or pathology associated with NOVX, treating a pathological state in a mammal, a vector comprising the activity of NOVX, treating or preventing a pathology associated with NOVX, treating a pathological state in a mammal, a vector comprising the NOVX nucleic acid molecule, a cell comprising the vector, an antibody that immunospecifically binds to NOVX, determining the presence or amount of NOVX or the nucleic acid molecule in a first mammalian subject. The methods and compositions of the present invention are useful for the diagnosis and creation of disorders associated with aberrant expression or activity of treatment of disorders associated with aberrant expression or activity of treatment of activity of the diagnosis and treatment of disorders associated with aberrant expression or activity of treatment.
                                                                                                                                                                                                                                                                                                                                    the NOVX polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and infilammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing, gene therapy and predictive medicine. The present sequence represents a NOVX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; NOVX; secreted protein; cancer; diabetes; obesity; endocrine disorder; CNS disorder; inflammatory disorder; gene therapy.
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100.0%; Pred. No. 2.3
:ive 0; Mismatches
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04-JUN-2002; 2002US-0385504P.
06-JUN-2002; 2002US-0386451P.
06-JUN-2002; 2002US-0386453P.
06-JUN-2002; 2002US-0386974P.
07-JUN-2002; 2002US-0386976P.
07-JUN-2002; 2002US-0387540P.
11-JUN-2002; 2002US-0387559P.
12-JUN-2002; 2002US-0387559P.
12-JUN-2002; 2002US-03875540P.
13-JUN-2002; 2002US-03875540P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted protein NOV2c.
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2002US-0389742P.
2002US-0390006P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 401 AA;
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17-JUN-2002;
19-JUN-2002;
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Zhong M;
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lo X, Gusev VY, Ji W;
Ort T, Padigaru M;
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                                                                                                                                                                                                                                                                                                                     Human; NOVX; secreted protein; cancer; diabetes; obesity; endocrine disorder; CNS disorder; inflammatory disorder; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a new isolated polypeptide (designated NOVX) comprising one of 141 fully defined sequences, their mature forms, a
                                                       Gaps
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Morman L, Guo A, Padigaru

JM, Millet I, Ort T, Padigaru

CEA, Peyman JA, Rieger DK;

SG, Smithson G, Spytek KA, F
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                        DB 8; Length 401;
                                       2.3e+02;
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A A, Ellerman K, Gerlach VL, Gorman.
A Macdougall JR, Malyankar UW, Mille
A SK, Patturajan M, Pena CEA, Peyras SK, Sciore P, Shenoy SG, Smi
                              100.0%; Preq. ....
         30.0%; Score 6; D
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                                                                                                                                                                                             ADJ34123 standard; protein; 401 AA.
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2002US - 03868 L6P.
2002US - 0387540P.
2002US - 0387559P.
2002US - 0387659P.
2002US - 0387659P.
2002US - 0389723P.
2002US - 0389729P.
2002US - 0389729P.
                                                                                                                                                                                                                                                                                         Human secreted protein NOV2d.
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2002US-0403748P.
2002US-0387037P.
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2002US-0386453P.
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2002US-0402832P.
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                                                                                                                                                                                                                                                           (first entry)
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Rothenberg ME, Sciore P,
Taupier RJ, Tchernev VT,
                                                     6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
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N-PSDB; ADJ34122.
                                     Local Similarity
                                                                                                                60 WSSWAL 65
                                                                                    3 WSSWAL 8
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07-JUN-2002;
10-JUN-2002;
11-JUN-2002;
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17-JUN-2002; 2
19-JUN-2002; 2
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                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                          Query Match
                                        Best Loc
Matches
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                                                                                                                                                                                ADJ34123
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Gaps

Human secreted protein NOV2f.

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The invention relates to a new isolated polypeptide (designated NOVX) comprising one of 141 fully defined sequences, their mature forms, a protein comprising one or more conservative substitutions or having at least 95% identity to one of the 141 proteins. Also included are a composition comprising NOVX (or a NOVX nucleic acid molecule (NA)), a kit composition to composition of NOVX incleic acid molecule encoding a NOVX protein, producing NOVX (comprising prolypeptide, where the call comprises a vector comprising NOVX (comprising polypeptide, where the call comprises a vector comprising NOVX (supering therefore agent that binds to NOVX, identifying a potential therapeutic agent for use in the treatment of a pathology that is related to abstrant expression or physiological interactions of NOVX, screening for a modulator of activity of or latency or predisposition to a pathology associated with NOVX, modulating the activity of NOVX, treating cor preventing a pathology associated with NOVX, unaminating the pathology associated with NOVX, treating a pathology associated with NOVX, treating call comprising the vector, an antibody that immunospecifically binds to molecule in a sample. And determining the presence or amount of NOVX, treating correction and persence or amount of NOVX or hundred or or preventing and determining the presence or amount of NOVX or hundred or or preventing and determining the presence or amount of NOVX or hundred or or preventing and determining the molecule in a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        compositions of the present invention are useful for the diagnosis and treatment of disorders associated with aberrant expression or activity of the NOVY polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and inflammatory disorders. They can be be used in various detection and screening assays, chromosome mapping, tissue typing, gene therapy and predictive medicine. The present sequence represents a NOVX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        molecule in a sample, and determining the presence of or predisposition to a disease associated with altered levels of expression of NOVX or the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated NOVX polypeptides useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory or CNS diseases.
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rman L, Guo X, Gusev VY, Ji
Millet I, Ort T, Padigaru M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid molecule in a first mammalian subject. The methods and
                                                                                                                                                                                                      , Burgess CE, Casm., Guo ...
Gerlach VL, Gorman L, Guo ...
Malyankar UM, Millet I, Ort T, roll
Pena CEA, Peyman JA, Rieger DK;
TSG, Smithson G, Spytek KA, S
                                                                                                                                                                                                         DW, Boldog FL, Burge..., Gorman, Macdougall JR, Malyankar UM, Mill', SK, Patturajan M, Pena CEA, Pey T, Sciore P, Shenoy SG, Smire. Sciore P, Shenoy SG, Smire.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 10; 418pp; English.
                      , 2002US-0402832P.

; 2002US-0403486P.

; 2002US-0403522P.

; 2002US-0403748P.

; 2002US-0387037P.

; 2003US-00454246.
                                                                                                                                                                                                                                                                                   Macdougall JR, Maly
a SK, Patturajan M,
                                                                                                                                                                                          (CURA-) CURAGEN CORP.
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                                            13-AUG-2002;
                                                                                           15-AUG-2002;
06-NOV-2002;
                                                                                                                                           03-JUN-2003;
17-JUL-2002;
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Rothenberg M
                                                                                                                                                                                                                                                                                                                                                           Taupier RJ,
                                                                                                                                                                                                                                       Anderson DW,
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30.0%; Score 6; DB 8; Length 401; 100.0%; Pred. No. 2.3e+02; tive 0; Mismatches 0; Indels Conservative Local Similarity es 6; Conserv Query Match Best Local Matches

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Gaps

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60 WSSWAL 65 œ 3 WSSWAL g

ADJ34127 standard; protein; 401 AA. ADJ34127; RESULT 43 ADJ34127 SXXXX

(first entry)

06-MAY-2004

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Human, NOVX, secreted protein, cancer; diabetes; obesity;
endocrine disorder; CNS disorder; inflammatory disorder; gene therapy.
                                                                                                                 2002US-0386816P.
2002US-0387002P.
2002US-0387540P.
                                                                                                                                         2002US-0387934P.
2002US-0389123P.
2002US-0389729P.
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2002US-0396706P.
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2002US-0386974P
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13-AUG-2002; 2002US-0403486P.
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                                                                        04-JUN-2003; 2003WO-US017512
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                                                WO2004000997-A2
                                                                                   19-MAR-2002;
04-dun-2002;
06-dun-2002;
06-dun-2002;
06-dun-2002;
07-dun-2002;
10-dun-2002;
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12-JUN-2002;
13-JUN-2002;
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                                    Homo sapiens
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M,;
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Stone Zhong

Stone DJ; Zhong M; Casman SJ, Edinger SR; man L, Guo X, Gusev VY, Ji illet I, Ort T, Padigaru M; Peyman JA, Rieger DK; CURAGEN CORP.

In DW, Boldog FL, Burgess CE, Casman, Guo X, C.,

A, Ellerman K, Gerlach VL, Gorman L, Guo X, C.,

Macdougall JR, Malyankar UM, Millet I, Ort T, Padigar

Macdugall JR, Pena CBA, Peyman JA, Rieger DK,

Patturajan M, Pena CBA, Peyman JA, Rieger DK,

The CBA, Peyman JA, Saytek KA,

Patturajan M, Bena CBA, Peyman JA, Serhusen BD,

CAM, Voss EZ, Zerhusen BD, Rothenberg Mi Taupier RJ, Prayaga SK, Anderson DW, Eisen A,

(CURA-) CURAGEN CORP.

WPI; 2004-082483/08. N-PSDB; ADJ34126.

New isolated NOVX polypeptides useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory or CNS diseases.

Claim 1; SEQ ID NO 16; 418pp; English.

The invention relates to a new isolated polypeptide (designated NOVX) comprising one of 141 fully defined sequences, their mature forms, a protein comprising one or more conservative substitutions or having at least 95% identity to one of the 141 proteins. Also included are a composition comprising NOVX (or a NOVX nucleic acid molecule (NA)), a kit comprising the composition of NOVX in one or more containers, an isolated culturing a cell under conditions that lead to expression of the polypeptide, where the cell comprises a vector comprising NOVX NA), identifying an agent that binds to NOVX, identifying a potential therapeutic agent for use in the treatment of a pathology that is related to aberrant expression or physiological interactions of NOVX, screening for a modulator of activity of or latency or predisposition to a pathology associated with NOVX, modulating the activity of NOVX, treating or preventing a pathology associated with NOVX, nucleic acid molecule, a state in a mammal, a vector comprising the NOVX nucleic acid molecule, a cell comprising the vector, an antibody that immunospecifically binds to NOVX, determining the presence or amount of NOVX or the nucleic acid molecule in a sample, and determining the presence of or predisposition genetic disorders or other traits to assess biodiversity

responsible for

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                                                          of
to a disease associated with altered levels or expression or not of included and compositions of the present invention are useful for the diagnosis and treatment of disorders associated with aberrant expression or activity of the NOVX polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing, gene detection and screening assays, chromosome mapping, tissue typing, gene
                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g; gene mapping; gene therapy; forensic;
imaging; diagnostic; genetic disorder.
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                                                                                                                                                                                                                    30.0%; Score 6; DB 8
100.0%; Pred. No. 2.3
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human diagnostic protein #14206.
                                                                                                                                                                                                                                                                                                                                                                                                                      ABG14215 standard; protein; 426 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001; 2001WO-US008631
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23-AUG-2000; 2000US-00649167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; chromosome mapping;
food supplement; medical in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                          6; Conservative
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                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                    Sequence 401 AA;
                                                                                                                                                                                                                                                                                                3 WSSWAL
                                                                                                                                                                                                                                                                                                                                60 WSSWAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       biodiversity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-FEB-2002
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                                                                                                                                                  protein.
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The invention relates to Pseudomonas aeruginosa polypeptides and the polymucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, of the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant of P. aeruginosa drugs, as templates for recombinant components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences of Pseudomonas species using biochip technology Sequences ABO67826-ABO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but, was obtained in electronic format from USPTO at
                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
         and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
                                                                                                                                                                                                             Gaps
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                                                                                                                                                                          Length 426;
                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                          DB 4; Len . 2.4e+02;
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                                                                                                                                                               30.0%; Sco. 100.0%; Pred. No. 2...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Deloughery C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pseudomonas aeruginosa polypeptide #5507.
                                                                                                                                                                                                                                                                                                                                                                     ABO73332 standard; protein; 478 AA.
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98US-0094190P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                               6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pseudomonas aeruginosa.
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                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                 78 GWRWLR
                                                                                                                                         Sequence 426 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-FEB-1999;
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ID ABO7
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Sequence 478 AA;

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Local Similarity 100.
Les 6; Conservative
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                                                      Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                     LRRYGW 18
                                                                                                                                                                                                   N-PSDB; AAL43410.
                                                                                                                                                                                                                                                                                                                                                                             Sequence 502 AA;
        A thaliana GAD1
                                                                                                                                                                       Kinnersley AM,
                                                                         WO200238736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                            16-MAY-2002
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Domain
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to identifying target proteins (ABB90790-ABB94016) for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant organisms using suitable search parameters, where plant sequences having an E-value greater by a factor of 3 than the E-value of most similar non-plant sequences are selected. The polypeptides or nucleic acids encoding them are useful for identifying modulators. The identified modulators are useful as
                                                                                                                                                                                                                                                                                                                                                                  Identifying plant target proteins for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                           Gaps
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       DB 7; Length 478;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30.0%; Score 6; DB 5; Length 502; 100.0%; Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                          Indels
                                                                                                                                                                      Herbicidally active polypeptide SEQ ID NO 2732.
      30.0%; Score 6; DB 7
100.0%; Pred. No. 2.6
ive 0; Mismatches
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0; Mismatches
                                                                                                                                                                                          Herbicidal; plant; agriculture; herbicide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA015132 standard; protein; 502 AA
                                                                                                              ABB93521 standard; protein; 502 AA.
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                                                                                                                                                     (first entry)
                           6; Conservative
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                                                                                                                                                                                                                                                                                                                             Weidler M;
                                                                                                                                                                                                             Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                               WPI; 2002-269010/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              395 LRRYGW 400
                                                                443 WRWLRR 448
        Query Match
Best Local Similarity
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                                             10 WRWLRR 15
                                                                                                                                                                                                                                                                                                          (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 502 AA;
                                                                                                                                                                                                                               WO200210210-A2
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                          Matches
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The present invention relates to a method of producing a transformed plant that selectively increases production of gamma-aminobutyric acid (GABA) in response to a signal, by incorporating into the plant's genome a DNA construct with a non-constitutive promoter operably linked to a polynucleotide encoding a functional plant glutamic acid decarboxylase (GAD), to provide a transformed plant that expresses the GAD coding sequence in response to a signal. Plants of this type have an enhanced ability to tolerate environmental or other stresses. The present sequence is the A. thaliana GADI protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Making transformed plants that selectively increase gamma-aminobutyric acid production, by incorporating a DNA construct with a polynucleotide encoding a plant glutamic acid decarboxylase enzyme into plant's genome
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GAD; plant GABA production regulation; glutamic acid decarboxylase; plant stress; GABA; gamma-aminobutyric acid; stress resistance.
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100.0%; Pred. No. 2.7
:ive 0; Mismatches
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/label= Cytoplasmic_domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EMER-) EMERALD BIOAGRICULTURE CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 17; Page 53; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                  07-NOV-2001; 2001WO-US047447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-NOV-2000; 2000US-0246367P
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                                                                                                                                                                                                          The present sequence is a human OCTIP (organic cation transporter-like protein), a member of the superfamily of sugar and other transporter molecules that have 12 transmembrane domains. The sequence is derived from a human foetal brain cDNA library. The protein is highly expressed an brain tissue and has nootropic, neuroprotective, neuroleptic, anticonvulsant, antiParkinsonian, antidepressant activities. The present sequence is used to regulate a variety of cellular processes e.g. cell proliferation, differentiation and survival, screen octip modulators and detect mutation in OCTIP gene. OCTIP modulators can be used to treat or prevent chronic neurodegenerative disorders (e.g. Alzheimer's, Parkinson's , Huntington's and ALS), CNS disorders (e.g. Alzheimer's, panic, depression), behavioural, sleep and eating disorders
                                                                                                                New nucleic acid encoding human organic cation transporter-like protein, used for prevention, treatment and diagnosis of e.g. neurological,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pain disorder; inflammatory pain; chronic pain; neuropathic pain; causalgia; fibromyalgia; cancer pain; migraine; headache; tissue pain; cardiovascular disorder; analgesic; human; 57145; OCT-5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30.0%; Score 6; DB 3; Length 548; 100.0%; Pred. No. 2.9e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                  Claim 9; Fig 1; 100pp; English.
                                                                                                                                   used for prevention, treatment
behavioral or sleep disorders.
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 (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-2001; 2001US-0333073P.
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                                Glucksmann MA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-441438/41.
                                                                 WPI; 2000-137069/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    235 LGWRWL 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                                                                 N-PSDB; AAZ49686.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 548 AA;
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                                  Goodearl AJ,
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71. 524
/note= "shows homology to a consensus sequence for sugar
and other transporter molecules derived from a hidden
Markov model"
                                                                                                                                                                                                                                                                176. .181
/label= Extracellular_domain
/note= "Hydrophilic region useful for antibody
                                                                                                                                                                                                                                                                                                                                                                                                                               233. .238
/label= Extracellular_domain
/note= "Hydrophilic region useful for antibody
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|Tabel= Extracellular domain
'note= "Hydrophilic region useful for antibody
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/label= Extracellular domain
prode= "Hydrophilic region useful for antibody
production"
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|Jabel= Extracellular_domain
|note= "Hydrophilic region useful for antibody
                                                                                                                                note= "Hydrophilic region useful for antibody
                                                                                           109 .121
| Jabel = Extracellular_domain
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|label= Cytoplasmic_domain
|155. .175
|label= Transmembrane_domain
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|label= Cytoplasmic_domain
58. .479
|label= Transmembrane_domain
                                                                                                                                                                  .22. .140
label= Transmembrane_domain
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/label= Transmembrane_domain
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label= Transmembrane_domain
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/label= Transmembrane_domain
                                                                               'label= Transmembrane_domain
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label= Transmembrane_domain
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'label= Cytoplasmic_domain
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/label= Cytoplasmic_domain
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/label= Cy
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|Jabel= Tr
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disease, Alzheimer's

Tourette's

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Gaps

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0; Indels

8; Length 548;

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depression, anxiety, cerebral ischaemia, myotonia, stroke, Tourette's syndrome, neonatal cerebral haemorrhage, Parkinson's disease, Alzheim disease and dementia. The present sequence represents the amino acid sequence of the rat synaptic vesicle protein 2, SVOP.
                                                                                                          an 30.0%; Score 6; DB 8; Le. Similarity 100.0%; Pred. No. 2.9e+02; 6; Conservative 0; Mismatches 0;
                                                                                                                        Best Local Similarity
                                                                             Sequence 548 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                             AD077827;
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                                                                                                                                                                                                                                                 RESULT
   8X33333
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                                                                                                                                                                                                                                                                                The present invention relates to a method of treating a subject having a pain disorder characterised by aberrant 577, 20739 or 57145 polypeptide activity or nucleic acid expression. The method involves administering a modulator of 577 (also known as sodium dependent proline transporter). 20739 (also known as P21-activated kinase 3 (PpK-3)) or 57145 (also known as OCT-5). The 577, 20739 or 57045 modulator is useful for treating pain disorders like inflammatory pain, chronic pain, neuropathic pain, causalgia, fibromyalgia, cancer pain, migraine/headache pain and tissue pain. The invention also describes methods for diagnosis and prognosis of various cardiovascular disorders, and for identification of subjects
                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          convulsion; depression; anxiety; cerebral ischaemia; myotonia; stroke;
Tourette's syndrome; neonatal cerebral haemorrhage; Parkinson's disease;
Alzheimer's disease; dementia; synaptic vesicle protein 2; SVOP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of identifying a binding partner for synaptic vesicle protein 2 (SV2), involves incubating SV2 protein with levetiracetam and a potential binding partner and determining if the potential binding partner modulates the binding of levetiracetam to the SV2 protein or fragment, thus identifying a binding partner for the SV2 protein. The method is useful for identifying a binding partner for synaptic vesicle protein 2 and the identified binding partner for synaptic vesicle protein 2 and the identified binding partner is useful for treating epilepsy, epileptogenesis, seizure disorders, convulsions,
                                                                                                                                                                                 exhibiting a predisposition to such conditions. The present sequence is human 57145 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying binding partner for synaptic vesicle protein 2 (SV2) for treating epilepsy, Parkinson's disease, Alzheimer's disease, involves determining binding partner which modulates binding of levetiracetam o
                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rat synaptic vesicle protein 2, SVOP
                                                                                                                                                                                                                                                                                                                                                                                                                                   AD077835 standard; protein; 548 AA.
Claim 1; Page 83-84; 84pp; English
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                                                                                                                                                                                                                                                                          Best Local Similarity
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                                                                                                                                                                                                                                Sequence 548 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; levetiracetam; epilepsy; epileptogenesis; seizure disorder;
convulsion; depression; anxiety; cerebral ischaemia; myotonia; stroke;
Tourette's syndrome; neonatal cerebral haemorrhage; Parkinson's disease;
Alzheimer's disease; dementia; synaptic vesicle protein 2; SVOP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying binding partner for synaptic vesicle protein 2 (SV2) for treating epilepsy, Parkinson's disease, Alzheimer's disease, involves determining binding partner which modulates binding of levetiracetam or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human synaptic vesicle protein 2, SVOP, DNA.
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100.0%; Pred. No. 2.9
:ive 0; Mismatches
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                                                                                                                                                                                                                                                           AD077827 standard; protein; 548 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      its analog to SV2 protein.
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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8 LGWRWL 13
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Best Local Similarity
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                                                                         235 LGWRWL
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Mismatches

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6; Conservative

Matches

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The invention relates to a novel method for treating a neurological disorder associated with synaptic vesicle function, endocrinopathy or hormonal disease which comprises administering a compound or agent that modulates a function or activity of an SV2 (synaptic vesicle protein 2) protein. The method of the invention has anticonvulsant, neuroleptic, antimantc, antidepressant, tranquiliser, antimigraine, analgesic, cerebroprotective, vasotropic, vulnerary, muscle relaxant, neoropic, cerebroprotective, vasotropic, vulnerary, muscle relaxant, noctropic, relaxant, haemostatic, neuroprotective, antiparkinsonian, noctropic, relaxant, na antidiabetic applications. The method may be useful for treating a neurological disorder, endocrinopathy or hormonal disease associated with synaptic vesicle function. The neurological disorder is cosmitive disorders, movement disorders, bipolar disorders's disease, cognitive disease and dementia. The endocrinological disorder is chosen from endocrinopathies involving hypersecretion or hyposecretion of at least one hormone, gigantism, dwarfism, actmal-medulla-related diseases, hypoglycaemia and circulation shock. The current sequence is that of the human synaptic vesicle protein OP (SVOP) of the invention.
                                                                                                                                                                                                                                                                                                          neurological disorder; synaptic vesicle function; endocrinopathy; hormonal; SV2; synaptic vesicle protein 2; anticonvulsant; neuroleptic; antimimant; antidepressant; tranquiliser; antimigraine; analgesic; cerebroprotective; vasotropic; vulnerary; muscle relaxant; acaricide; relaxant; haemostatic; neuroprotective; antiparkinsonian; nootropic; endocrine; enidabetic; seizure; epilepsy; Parkinson's; Alzheimer's; cognitive; movement; bipolar; neurodegenerative; dementia; gigantism; dwarfism; adrenal-medulla; hypoglycaemia; circulation shock; human; SVOP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating neurological disorder associated with synaptic vesicle function such as seizure, epilepsy, endocrinopathy or hormonal diseases, involves administering compound that modulates function or activity of SV2
                                                                                                                                                                                                                                                                     Human synaptic vesicle protein OP (SVOP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 8; 135pp; English.
                                                                                                                                            ADP44588 standard; protein; 548 AA.
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30-SEP-2003; 2003US-0506764P.
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                                                                                                                                                                                                                                (first entry)
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N-PSDB; ADP44587.
                                       235 LGWRWL 240
LGWRWL 13
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                                                                                                                                                                                      ADP44588;
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The invention relates to a novel method for treating a neurological disorder associated with synaptic vesicle function, endocrinopathy or hormonal disease which comprises administering a compound or agent that modulates a function or activity of an SV2 (synaptic vesicle protein 1) protein. The method of the invention has anticonvulsant, neuroleptic, antidepressant, tranquiliser, antimigraine, analgesic, cerebroprotective, vasotropic, vulnerary, muscle relaxant, acaricide, relaxant, haemostatic, neuroprotective, antiparkinsonian, nootropic, relaxant, haemostatic, neuroprotective, antiparkinsonian, nootropic, reating a neurological disorder, endocrinopathy or hormonal disease associated with synaptic vesicle function. The neurological disorder is chosen from seizure, epilepsy, Parkinson's disease, Alzheimer's disease, neurodegenerative disease and demenia. The endocrinological disorder is chosen from endocrinopathies involving hypersecretion or hyposecretion of at least one hormone, gigantism, dwarfism, admenal-medulla-related diseases, hypoglycaemia and circulation shock. The current sequence is that of the Norway rat synaptic vesicle protein OP (SVOP) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating neurological disorder associated with synaptic vesicle function such as seizure, epilepsy, endocrinopathy or hormonal diseases, involves administering compound that modulates function or activity of SV2
                                                                                                                                                                                                                                                                                                                                                                                                       endocrine; antidiabetic; seizure; epilepsy; Parkinson's; Alzheimer's; cognitive; movement; bipolar; neurodegenerative; dementia; gigantism; dwarfism; adrenal-medulla; hypoglycaemia; circulation shock; Norway rat;
                                                                                                                                                                                                                                                                                                  neurological disorder, synaptic vesicle function, endocrinopathy, hormonal; SV2, synaptic vesicle protein 2; anticonvulsant; neuroleptic; antimanic; antidepressant; tranquiliser; antimigraine, analgesic; cerebroprotective; vasotropic; vulnerary; muscle relaxant; acaricide; relaxant; haemostatic; neuroprotective, antiparkinsonian; nootropic;
                                                                                                                                                                                                                                                                  Norway rat synaptic vesicle protein OP (SVOP).
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                                                                                                                                          ADP44596 standard; protein; 548 AA.
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30-SEP-2003; 2003US-0506764P.
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                                        235 LGWRWL 240
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8 LGWRWL 13
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                                                                                                  RESULT 53
                                                                                                                       ADP44596
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Sequence 548 AA;

Length 548;

Score 6; DB 8; Le Pred. No. 2.9e+02;

30.0%; S 100.0%;

Query Match Best Local Similarity

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Gaps

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enantiomeric conversion; amine; homochiral; enzyme; oxidation catalysis; stereoselective; microbial; monoamine oxidase; MAO; enantioselectlvity; deracemization; mutant; mutein.
                                                                                                                                                                                                                                                                                                                              /note= "The wild-type residue of Asn is substituted with
                                                                                                                                                                                                                                                                                                                                                            /note= "The wild-type residue of Met is substituted with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enantiomeric conversion of amines comprises treating a homochiral amine or mixture of amine enantiomers with enzyme capable of catalyzing oxidation of amine and subsequently or simultaneously treating with a
                                                                                                                                                                                    Aspergillus niger monoamine oxidase protein variant N336S + M348K.
                       0;
                                                                                                                                                                                                                                                                                           Misc-difference 1. .610 /note= "All Xaa residues signify a stop codon"
 DB 8; Length 548; 2.9e+02;
                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mahmoudian M;
30.0%; Score 6; DB 8
100.0%; Pred. No. 2.9
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Turner NJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 26; SEQ ID NO 3; 43pp; English.
                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                     ADD15932 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-2003; 2003WO-GB001198.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-MAR-2002; 2002GB-00006415.
                                                                                                                                                               (first entry)
                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alexeeva MV, Enright A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                                                                   348
                                                               235 LGWRWL 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-779267/73
          Local Similarity
                                           8 LGWRWL 13
                                                                                                                                                                                                                                                  Synthetic.
Aspergillus niger.
                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                              WO2003080855-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reducing agent,
                     9
                                                                                                                                                               15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                   02-OCT-2003
                                                                                                                                          ADD15932;
 Query Match
            Best Loc
Matches
                                                                                              RESULT 54
                                                                                                          ADD15932
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel method for the enantiomeric conversion of amines. The method comprises treating a homochiral amine or a mixture of amine enantioners with an enzyme capable of catalyshing oxidation of the amine in a stereoselective manner and, subsequently or simultaneously, treating with a reducing agent. A further method of the invention is useful for directing the evolution of an amine oxidation enzyme e.g., microbial monoamine oxidase (MAO), preferably Aspergillus niger monoamine oxidase or its variant. This method is also useful for increasing the enantioselectivity of an amine oxidase enzyme. The enantioselective amine oxidase enzyme of the invention is useful in deracemization of amines. This sequence represents a variant Aspergillus niger monoamine oxidase protein of the invention.
                                                                                                                                                                                                                                                                                                                      catalysis;
                                                                                                                                                                                                                                                                                                                 enantiomeric conversion; amine; homochiral; enzyme; oxidation catalysis stereoselective; microbial; monoamine oxidase; MAO; enantioselectivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enantiomeric conversion of amines comprises treating a homochiral amine or mixture of amine enantiomers with enzyme capable of catalyzing oxidation of amine and subsequently or simultaneously treating with a
                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "All Xaa residues signify a stop codon"
, DB 7; Let.5
MA, 3.2e+02;
0; Indels
                                                                                                                                                                                                                                                                                   Aspergillus niger monoamine oxidase protein variant N336S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30.0%; Score 6; DB 7; Length 610; 100.0%; Pred. No. 3.2e+02; live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Turner NJ, Mahmoudian M;
             30.0%; Score 6; DB 7
100.0%; Pred. No. 3.2
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 24; SEQ ID NO 2; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          key Location/Qualifiers Misc-difference j. .610
                                                                                                                                                                                          ADD15931 standard; protein; 610 AA.
                                                                                                                                                                                                                                                                                                                                                  deracemization; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-2003; 2003WO-GB001198.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-MAR-2002; 2002GB-00006415
                                                                                                                                                                                                                                                     (first entry)
Query Match
Best Local Similarity luv.
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Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Enright A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-779267/73.
                                                                                                           WALGWR 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference 336
                                                                             WALGWR 11
                                                                                                                                                                                                                                                                                                                                                                                                Aspergillus niger.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 610 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003080855-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reducing agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alexeeva MV,
                                                                                                                                                                                                                                                     15-JAN-2004
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                                                                                                                                                                                                                        ADD15931;
                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                             9
                                                                                                            459
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                                                                                                                                                           RESULT 55
                                                                                                                                                                           ADD15931
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the amine in a stereoselective manner and, subsequently or simultaneously, treating with a reducing agent. A further method of the invention is useful for directing the evolution of an amine oxidation enzyme e.g., microbial monoamine oxidase (MAO), preferably Aspergillus niger monoamine oxidase or its variant. This method is also useful for increasing the enantioselectivity of an amine oxidase enzyme. The enantioselective amine oxidase enzyme of the invention is useful in deracemization of amines. This sequence represents a variant Aspergillus niger monoamine oxidase protein of the invention.

Sequence 610 AA;

The invention relates to a novel method for the enantiomeric conversion of amines. The method comprises treating a homochiral amine or a mixture of amine enantiomers with an enzyme capable of catalysing oxidation of

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g à

ADD15930;

RESULT 56 ADD15930

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infection, and in detection of P. aeruginosa sequences or other sequences of Beeudomonas species using biochip technology Sequences ABO67826-
BOB64396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P aeruginosa drugs, as templates for recombinant production of P. aeruginosa drugs, as templates for recombinant production of P. aeruginosa drugs, as templates for accombinant production of P. aeruginosa-caused
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                predictor set; protein tyrosine kinase activity modulator; protein tyrosine kinase; cytostatic;
                                                                                                                                                     Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 7; Length 800; 3.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human src biomarker polypeptide SEQ ID NO:339.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 24635; 455pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30.0%; Score 6; DB 7
100.0%; Pred. No. 3.9
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nolling J, Deloughery C,
                                                                                                           Pseudomonas aeruginosa polypeptide #8064.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD14150 standard; protein; 880 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seqdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                             98US-0074788P.
98US-0094190P.
                                                                                                                                                                                                                                                                                                                                     99US-00252991.
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                                                               29-JUL-2004 (first entry)
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Best Local Similarity 100...
6; Conservative
                                                                                                                                                                                                    Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-615309/58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; ABD09460.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rubenfield MJ,
                                                                                                                                                                                                                                                                                                                                     18-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JAN-2004
                                                                                                                                                                                                                                                 US6551795-B1
                                                                                                                                                                                                                                                                                                                                                                                  18-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                    27-JUL-1998;
                                                                                                                                                                                                                                                                                           22-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD14150;
                     ABO75889;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD14150
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                                                                                                                                                                                                                                                                                                                                   enantiomeric conversion; amine; homochiral; enzyme; oxidation catalysis; stereoselective; microbial; monoamine oxidase; MAO; enantioselectivity; deracemization; wild-type.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enantiomeric conversion of amines comprises treating a homochiral amine or mixture of amine enantiomers with enzyme capable of catalyzing oxidation of amine and subsequently or simultaneously treating with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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/note= "All Xaa residues signify a stop codon"
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                                                                                                                                                                                                                                                                                             Wild-type Aspergillus niger monoamine oxidase protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mahmoudian M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7; SEQ ID NO 1; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Turner NJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                            ADD15930 standard; protein; 610 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-MAR-2003; 2003WO-GB001198
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                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alexeeva MV, Enright A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-779267/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WALGWR 464
                                              459 WALGWR 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6 WALGWR 11
    11
                                                                                                                                                                                                                                                                                                                                                                                                                              Aspergillus niger.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 610 AA;
    WALGWR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003080855-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reducing agent
                                                                                                                                                                                                                                                   15-JAN-2004
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Ď, Bush

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Gaps

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0; Indels

ABO75889 standard; protein; 800 AA.

RESULT 57 ABO75889 ID ABO75

459

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The present invention describes a predictor set comprising a plurality of polynucleotides or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein tyrosine kinase activity or members of the protein tyrosine kinase control of pathway. Also described: (1) predicting whether a compound is capable of modulating the activity of cells, comprising obtaining whether the cells, comprising botanining whether the cells, (2) a plurality of markers, and certaining the expression of the markers to the compound's ability to correlate which compound sensitivity or resistance of cells associated with a disease state; and (3) identifying polymucleotides and polypeptides where so not or more compounds sensitivity or resistance of cells associated with a disease state, comprising the expression pattern of a microarray of polymucleotides or polypeptides, and selecting the plurality or cell lines to one or more compounds, analysing the expression pattern of cells associated with a disease state by using the compounds that interact with polymucleotides and polypeptides are useful in predicting the ceststance of cells associated with a disease state by using the ceststance of cells associated with a disease state by using the ceststance of cells associated with a disease state by using the ceststance of cells associated with a disease state by using the compounds that interact with protein tyrosine kinase pathways. These may be used in determining drug centainity of compounds that interact with protein tyrosine kinase pathways. These may be used in determining deneric profiles which aid in treating diseases and disorders (e.g. cancer) based on patient response at a molecular level. The present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
  gene therapy; drug sensitivity; genetic profile; cancer; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human EST encoded protein SEQ ID NO: 1775.
                                                                                                                                                                                                                                                                                                        Shaw P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 10; SEQ ID NO 339; 139pp; English.
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                                                                                                                                                                                                                                                                                                        Lee FY,
                                                                                                                                                                                                                                                                 ဥ
                                                                                                                                                                       17-JAN-2003; 2003WO-US001981.
                                                                                                                                                                                                                    18-JAN-2002; 2002US-0350061P.
                                                                                                                                                                                                                                                             (BRIM ) BRISTOL-MYERS SQUIBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                        Huang F, Fairchild CR,
                                                                                                                                                                                                                                                                                                                                                WPI; 2003-636735/60.
N-PSDB; ADD14752.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 880 AA;
                                                                                     WO2003062395-A2
                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-OCT-2001
                                                                                                                              31-JUL-2003
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AAM24250
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BXHXHXB
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             Human, sheep, pig, cow, fruit fly, yeast, hamster, macaque, horse,
tomato, monkey, dog, sea urchin, expressed sequence tag, EST,
diagnostics, forensic test, gene mapping, genetic disorder, biodiversity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (ESTs) from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, semaphorin G, NHP, semaphorin homologue, therapeutic, diagnostic, pharmacogenomic, drug screening, clinical trial, physiological disorder, physiological disease, cosmetic, nutriceutical.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                Isolated polypeptide for treatment of diseases, diagnostics, raising antibodies and research use.
                                                                                                                                                                                                                                                                                                              Asundi V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 896;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                              Chen R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human semaphorin G-like NHP protein SEQ ID NO:16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4; Len
                                                                                                                                                                                                                                                                                                           ou P, Qian XB, Wang Z,
Zhang J, Werhman T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30.0%; Score 6; DB 4
100.0%; Pred. No. 4.3
live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 20; Page 1166-1168; 1275pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG68296 standard; protein; 939 AA.
                                                                                                                                                                        25-JAN-2001; 2001WO-US002687.
                                                                                                                                                                                                    25-JAN-2000; 2000US-00491404.
17-UUD-2000; 2000US-00617746.
03-AUG-2000; 2000US-00631451.
15-SEP-2000; 2000US-00663870.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-MAY-2001; 2001WO-US015489.
                                                                                                                                                                                                                                                                                                           Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-FEB-2002 (first entry)
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Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the invention
                                                          gene therapy; nutrition.
                                                                                                                                                                                                                                                                                                           Tang YT, Liu C, Zho
Cao Y, Drmanac RA,
                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-476164/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         687 WLRRYG 692
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                                                                                                                                                                                                                                                                             (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; AAH98909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 896 AA;
                                                                                                                 WO200154477-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200188133-A2.
                                                                                     Homo sapiens
                                                                                                                                             02-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAG68296;
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AAG68296
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Turner CA;

Scoville J,

Wang X,

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New nucleic acid molecule that encodes human semaphorin homolog, usef
in therapeutic, diagnostic and pharmacogenomic applications, in drug
screening and in clinical trial monitoring.
02-JUN-2000; 2000US-0208893P.
                            (LEXI-) LEXICON GENETICS INC.
                                                                              WPI; 2002-055698/07.
N-PSDB; ABA04068.
                                                      Walke DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY94990
    à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             g
                                                                                                                                                                                                                                                                 in therapeutic diagnostic and pharmacogenomic applications, in drug screening, clinical trial monitoring, the treatment of physiological disorder or diseases, and in cosmetic or mutriceutical applications. (I) can be used for identifying coding sequences and mapping a unique gene to can be used for identifying coding sequences and mapping a unique gene to prepare cloning and sequencing templates, as hybridisation probes for screening libraries and assessing gene expression patterns, in microarrays and other assay formats to screen collections of genetic material from patients who have a particular medical condition, to compare in a particular medical condition, to controprose that are useful for gene requlation. (I) can also be used for the diagnostic and/or triple helix sequences that are useful for gene requlation. (I) can also be used for the diagnosis of disease, for screening for drugs effective in the treatment of symptomatic or phenotypic manifestations of perturbing the normal function of a human semaphorin homologue in the last are at at a service and by the articular midgenesis/evolution of proteins that are at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ò
                                                                                                                                                                                                                                ABA04062 to ABA04087 represent polynucleotide sequences (I) which encode the human semaphorin homologous proteins (particularly similar to semaphorin G) given in AAG68289 to AAG68312 (II). (I) and (II) are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; semaphorin G; NHP; semaphorin homologue; therapeutic; diagnostic; pharmacogenomic; drug screening; clinical trial; physiological disease; cosmetic; nutriceutical.
                                                                                                                                                  New nucleic acid molecule that encodes human semaphorin homolog, useful in therapeutic, diagnostic and pharmacogenomic applications, in drug screening and in clinical trial monitoring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human semaphorin G-like NHP protein SEQ ID NO:14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30.0%; Score 6; DB 5
100.0%; Pred. No. 4.5
iive 0; Mismatches
                                                                                  Turner CA;
                                                                                                                                                                                                        Claim 1; Page 62-64; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAG68295 standard; protein; 954 AA.
                                                                                  Scoville J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        partially encoded by (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-MAY-2001; 2001WO-US015489.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-MAY-2000; 2000US-0205274P.
                18-MAY-2000; 2000US-0205274P.
                                                         (LEXI-) LEXICON GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 1000.
                                                                                                             WPI; 2002-055698/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         473 WSSWAL 478
                                                                                    Wang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 939 AA;
                                                                                                                        N-PSDB; ABA04069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 WSSWAL
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                                                                                    Walke DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 61
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consistent of againstic and pharmacogenic applications, in drug screening, clinical trial monitoring, the treatment of physiological disorder or diseases, and in cosmetic or nutriceutical applications. (I) can be used for identifying coding sequences and mapping a unique gene to a particular chromosome, to screen libraries, to isolate clones, to prepare cloning and sequencing templates, as phyridistation probes for screening libraries and assessing gene expression patterns, in microarrays and other assay formats to screen collections of genetic microarrays and other assay formats to screen collections of genetic cidentify mutations associated with a particular disease, in a diagnostic or prognostic assay, as antisense molecule, and as part of ribozyme or prognostic assay, as antisense molecule, and as part of ribozyme corporations for the diagnostic state are useful for gene regulation. (I) can also be used for detecting mutant proteins or inappropriately corporate in the treatment of symptomatic or phenotypic manifestations of perturbing the normal function of a human semaphorin homologue in the coded by (I) least partially encoded by (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                      ABA04062 to ABA04087 represent polynucleotide sequences (I) which encode the human semaphorin homologous proteins (particularly similar to semaphorin G) given in AAG68289 to AAG68312 (II). (I) and (II) are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; secreted protein; cancer; tumour; cardiovascular disorder;
blood disorder; haemophilia; autoimmune disease; diabetes; inflammation;
infection; fungal; bacterial; viral; HIV; allergy; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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4.5e+02;
7.7 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neurodegenerative disease; asthma; contraceptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted protein vb21_1, SEQ ID NO:20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. No. 4.5 ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 6;
Claim 1; Page 59-61; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY94990 standard; protein; 999 AA.
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98US-0097659P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US019351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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24-AUG-1998;
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The invention relates to 40 human secreted proteins (AAY94981-Y95020),
and cDNA sequences encoding them (AAA23423-A23462). The secreted proteins
CC frie invention include those that are thought to be only partially
secreted, i.e., transmembrane proteins. The proteins of the invention may
cactivity; cell proliferation; differentiation; immune modulation;
cactivity; chemotactic/chemokinetic activity; nactivin/inhibin
activity; chemotactic/chemokinetic activity; haemostatic and thrombolytic
activity; chemotactic/chemokinetic activity; haemostatic and thrombolytic
activity; chemotactic/chemokinetic activity; and tumour inhibition activity. The
cc activity; anti-inflammatory activity; and tumour inhibition activity. The
proteins may be administered to patients as vaccines, and the nucleotides
cmay be treated using the proteins or nucleotides of the invention include
autoimmune diseases; genetic disorders; haemophila; cardiovascular
diseases; cancer; bacterial, tungal and viral infections, especially HIV;
cdiseases; cancer; bacterial, tungal and viral infeations, and
allergic reactions such as asthma and anaemia. They may also be used for
treating wounds; burns, ulcers, osteoporosis, osteoarthitis; pulmington's disease
cand amyotrophic lateral sclerosis (ALS). Proteins with activin/inhibin
cc activity may additionally be useful as contraceptives. Nucleic acid
cactivity may additionally be useful as contraceptives. Nucleic acid
cactivity may additionally be useful as contraceptives. Nucleic acid
cactivity may additionally be useful as contraceptives. Nucleic acid
cone of the 40 proteins of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ò
                                                                                                                                                                                                                                                   New secreted or transmembrane proteins and polynucleotides encoding them, useful for treating neurodegenerative disorders, autoimmune diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 30.0%; Score 6; DB 3; Length 999; Similarity 100.0%; Pred. No. 4.7e+02; 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                           Rapiejko P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human semaphorin G-like NHP protein SEQ ID NO:6.
                                                                                                                                                                         Hall J,
                                                                                                                                                                                                                                                                                                                           Claim 29; Page 276-280; 357pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAG68291 standard; protein; 1034 AA.
                                                                                                                                                                         Hoffman H,
                 98US-0102092P.
98US-0109978P.
98US-0113645P.
98US-0113646P.
 98US-0099618P
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                                                                                                                                                                         Yuan O,
                                                                                                                                   (ALPH-) ALPHAGENE INC
                                                                                                                                                                                                               WPI; 2000-224657/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     518 WSSWAL 523
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 WSSWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 999 AA;
                                                                                                                                                                       Valenzuela D,
                                     25-NOV-1998;
                                                         23-DEC-1998;
                                                                                             23-AUG-1999;
09-SEP-1998;
                                                                           23-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAG68291;
                                                                                                                                                                                                                                                                                           cancer.
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in therapeutic, diagnostic and pharmacogenomic applications, in drug screening, clinical trial monitoring, the treatment of physiological disorder or diseases, and in codming the treatment of physiological can be used for identifying coding sequences and mapping a unique gene to prepare cloning accounting templates, to isolate clones, to prepare cloning and sequencing templates, as hybridisation probes for screening libraries and assessing gene expression patterns, in microarrays and other assay formats to screen collections of genetic incoarrays and other assay formats to screen collections of genetic identify mutations associated with a particular medical condition, to identify mutations associated with a particular disease, in a diagnostic and/or triple helix sequences that are useful for gene regulation. (I) can also be used for detecting mutant proteins or inappropriately expressed proteins for the diagnosis of disease, for screening for drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    effective in the treatment of symptomatic or phenotypic manifestations of perturbing the normal function of a human semaphorin homologue in the body, and in the molecular mutagenesis/evolution of proteins that are at least partially encoded by (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA04062 to ABA04087 represent polynucleotide sequences (I) which encode the human semaphorin homologous proteins (particularly similar to semaphorin G) given in AAG68289 to AAG68312 (II). (I) and (II) are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, semaphorin G; MHP; semaphorin homologue; therapeutic; diagnostic; pharmacogenomic; drug screening; clinical trial; physiological disease; cosmetic; nutriceutical.
                                                                                                                                                                                                                                                                                                                                             New nucleic acid molecule that encodes human semaphorin homolog, usef in therapeutic, diagnostic and pharmacogenomic applications, in drug screening and in clinical trial monitoring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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. 4.8e+02;
-hag 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human semaphorin G-like NHP protein SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30.0%; Score 6; DB 5
100.0%; Pred. No. 4.8
iive 0; Mismatches
                                                                                                                                                                                                                                                      Turner CA;
/note= "encoded by RAC"
                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 44-47; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAG68289 standard; protein; 1049 AA
                                                                                                                                                                                                                                                    Walke DW, Wang X, Scoville J,
                                                                                                                                                  18-MAY-2000; 2000US-0205274P.
                                                                                                               14-MAY-2001; 2001WO-US015489
                                                                                                                                                                                                           (LEXI-) LEXICON GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Conservative
                                                                                                                                                                                                                                                                                        WPI; 2002-055698/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             568 WSSWAL 573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1034 AA;
                                                                                                                                                                                                                                                                                                             N-PSDB; ABA04064
                                    WO200188133-A2.
                                                                       22-NOV-2001
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Homo sapiens

Human, semaphorin G, NHP, semaphorin homologue, therapeutic, diagnostic, pharmacogenomic; drug screening, clinical trial, physiological disease, cosmetic, nutriceutical.

Key Location/Qualifiers Misc-difference 34

Homo sapiens

22-NOV-2001

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companyor to hardrane proteins ordering statements and the human semaphorin homologous proteins organizations in the human semaphorin homologous proteins (II). (I) and (II) are useful in therapeutic, diagnostic and pharma-openomic applications, in drug screening, clinical trial monitoring, extrement of physiological disorder or diseases, and in cosmetic or nutriceutical applications. (I) can be used for identifying coding sequences and mapping a unique gene to particular chromosome, to screen libraries, to isolate clomes, to prepare cloning and sequencing templates, as hybridisation probes for screening libraries and assessing gene expression patterns, in microarrays and other assay formats to screen collections of genetic material from patients who have a particular medical condition, to identify mutations associated with a particular disease, in a diagnostic or prognostic assay, as antisense molecule, and as part of ribozyme and/or triple helix sequences that are useful for gene regulation. (I) can also be used for detecting mutant proteins or inappropriately
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                can also be used for detecting mutant proteins or inappropriately expressed proteins for the diagnosis of disease, for screening for drugs effective in the treatment of symptomatic or phenotypic manifestations of perturbing the normal function of a human semaphorin homologue in the body, and in the molecular mutagenesis/evolution of proteins that are at least partially encoded by (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA04062 to ABA04087 represent polynucleotide sequences (I) which encode
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid molecule that encodes human semaphorin homolog, usef in therapeutic, diagnostic and pharmacogenomic applications, in drug screening and in clinical trial monitoring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, PRO polypeptide, membrane bound protein, receptor; diagnosis; transmembrane; secretion; immunoadhesion; pharmaceutical; screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Se+02;
5e+02;
7-1 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PR01249 (UNQ632) amino acid sequence SEQ ID NO:102.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30.0%; Score 6; DB 5
100.0%; Pred. No. 5e+
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Turner CA;
                                                                                                    /note= "encoded by RAC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 48-50; 103pp; English.
                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                   Scoville J,
                                                                                                                                                                                                                                                                                                       18-MAY-2000; 2000US-0205274P. 02-JUN-2000; 2000US-0208893P.
                                                                                                                                                                                                                                                     14-MAY-2001; 2001WO-US015489.
                                                                                                                                                                                                                                                                                                                                                                                 (LEXI-) LEXICON GENETICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002-055698/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang X,
                                                    Key
Misc-difference 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 WSSWAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1078 AA;
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N-PSDB; ABA04065.
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                                                                                                                                                      WO200188133-A2
     Homo sapiens.
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                                                                                                                                                                                                        22-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in therapeutic, diagnostic and pharmacogenomic applications, in drug screening, clinical trial monitoring, the treatment of physiological disorder or diseases, and in cosmetic or untriceutical applications. (I) can be used for identifying coding sequences and mapping a unique gene to a particular chromosome, to screen libraries, to isolate clones, to prepare cloning and sequencing templates, as hybridisation probes for screening libraries and assessing gene expression patterns, in microarrays and other assay formats to screen collections of genetic material from patients who have a particular medical condition, to identify mutations associated with a particular disease, in a diagnostic or prognostic assay, as antisense molecule, and as part of ribozyme and/or triple helix sequences that are useful for gene regulation. (I) can also be used for detecting mutant proteins or inappropriately
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expressed proteins for the diagnosis of disease, for screening for drugs effective in the treatment of symptomatic or phenotypic manifestations of perturbing the normal function of a human semaphorin homologue in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA04062 to ABA04087 represent polynucleotide sequences (I) which encode the human semaphorin homologous proteins (particularly similar to semaphorin G) given in AAG68289 to AAG68312 (II). (I) and (II) are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          perturbing the normal function of a human semaphorin homologue in the body, and in the molecular mutagenesis/evolution of proteins that are at least partially encoded by (I)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pharmacogenomic; drug screening; clinical trial; physiological disorder; physiological disease; cosmetic; nutriceutical.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid molecule that encodes human semaphorin homolog, useful in therapeutic, diagnostic and pharmacogenomic applications, in drug screening and in clinical trial monitoring.
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                                                                                                                                                                                                                                                                                                                                                                                    Turner CA;
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100.0%; Pred. No.
:ive 0; Mismatch
                                                       /note= "encoded by RAC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 37-39; 103pp; English.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                    Scoville J,
                                                                                                                                                                                                           14-MAY-2001; 2001WO-US015489.
                                                                                                                                                                                                                                                          18-MAY-2000; 2000US-0205274P, 02-JUN-2000; 2000US-0208893P.
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Best Local Similarity 100.
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                             Misc-difference 34
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Gaps

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AAG68292;

RESULT 65 AAG68292

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Human;

Length 1078;

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98US-0098716P

98US-0098742P

98US-0098831P

98US-0098831P

98US-0099836P

98US-0099536P

98US-0099536P

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98US-009963P

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98US-010234P

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98US-010234P

98US-010256P

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98US-010256P
                                                                   99WO-US020111
                           WO200012708-A2
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02-OCT-1998;
06-OCT-1998;
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17.5EP-1998
17.5EP-1998
18.5EP-1998
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18.5EP-1998
18.5EP-1998
18.5EP-1998
18.5EP-1998
22.5EP-1998
22.5EP-1998
         Homo sapiens
                                                                                             01-SEP-1998

02-SEP-1998

02-SEP-1998

03-SEP-1998

09-SEP-1998

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11-SEP-1998

11-SEP-1998
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23-SEP-1998;
24-SEP-1998;
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24-SEP-1998;
24-SEP-1998;
24-SEP-1998;
29-SEP-1998;
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29-SEP-1998,
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30-SEP-1998,
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30-SEP-1998,
30-SEP-1998,
                                                                 01-SEP-1999;
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17-SEP-1998;
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17-SEP-1998;
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23-SEP-1998;
23-SEP-1998;
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New mammalian DNA sequences encoding transmembrane, receptor or secreted PRO polypeptides, useful for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interactions.
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 98US-0103449P.
98US-0103314P.
98US-0103328P.
98US-0103339F.
98US-0103339F.
98US-0103633P.
98US-0103673P.
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N-PSDB; AAA37051.
06-OCT-1998)
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AAA37022 to AAA37144 encode the new isolated human transmembrane,

Claim 12; Fig 60; 773pp; English.

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Gaps

0;

DB 4; Le.\_, No. 5e+02; 0; Indels

30.0%; Score 6; DB 4 100.0%; Pred. No. 5e+ cive 0; Mismatches

Length 1089;

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6; Conservative
                                                                                                    687 WLRRYG 692
                                  Query Match
Best Local Similarity
                                                                               12 WLRRYG 17
             Sequence 1089 AA;
                                                       Matches
                                                                                                                                    RESULT 68
                                                                                                                                                 AAU29156
                                                                                                                                                                     XX OS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to secreted and transmembrane proteins. Hese proteins and the DNA encoding them may be used as hybridization probes, in chromosome and gene mapping and in the generation of antisense RNA and DNA. They may also be used used to generate either transgenic animals or knockout animals which are in turn useful for development and screening of therappeutically useful reagents. The nucleic acids may also be used in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hillan KJ;
Watanabe CK;
receptor or secreted PRO polypeptides given in AAY99340 to AAY99462. The transmembrane and receptor PRO proteins can be used for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interactions. The polypeptides and nucleotide sequences encoding then have various industrial applications, including uses as pharmaceutical and diagnostic agents. AAA37145 to AAA37330 represent PCR primers and hybridisation probes used in the isolation of the PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful as hybridization probes, in chromosome and gene mapping and gene
                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Secreted and transmembrane proteins and nucleic acids designated PRO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fong S;
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                                                                                                                          Length 1089;
                                                                                                                          30.0%; Score 6; DB 3; Length 1089
100.0%; Pred. No. 5e+02;
.ive 0; Mismatches 0; Indels
                                                                                 polypeptides from the present invention
                                                                                                                                                                                                                                                                                                                                         Secreted; transmembrane; gene therapy.
                                                                                                                                                                                                                                                  AAB66118 standard; protein; 1089 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0141037P.
99US-0144758P.
99US-0145698P.
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99WO-US030095,
2000WO-US000219,
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99US-0162506P.
                                                                                                                                                                                                                                                                                                                   Protein of the invention #30
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                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-071395/08
                                                                                                                                                                                           687 WLRRYG 692
                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                     12 WLRRYG 17
                                                                                                      Sequence 1089 AA;
                                                                                                                                                                                                                                                                                                                                                                                     WO200078961-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-SEP-1999;
29-OCT-1999;
30-NOV-1999;
02-DEC-1999;
16-DEC-1999;
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AAB66118
                                                                                                                                                 Matches
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PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep; dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha; blood; chondrocyte cell; cell proliferation; cell differentiation; colon; adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.
AAU29156 standard; protein; 1089 AA
                                                                                                                                                     Human PRO polypeptide sequence #133.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 MAR. 2000; 2000US-0189320P.

15-MAR. 2000; 2000US-0189320P.

15-MAR. 2000; 2000US-0189320P.

21-MAR. 2000; 2000US-01900884.

21-MAR. 2000; 2000US-0191007P.

21-MAR. 2000; 2000US-0191048P.

21-MAR. 2000; 2000US-0191314P.

28-MAR. 2000; 2000US-0193033P.

29-MAR. 2000; 2000US-0193033P.
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04-APR-2000; 2000W0-05094449P.
11-APR-2000; 2000WS-0194449P.
11-APR-2000; 2000WS-0195975P.
11-APR-2000; 2000WS-01960WP.
11-APR-2000; 2000WS-01960WP.
11-APR-2000; 2000WS-0196820P.
11-APR-2000; 2000WS-0198585P.
25-APR-2000; 2000WS-0199397P.
25-APR-2000; 2000WS-0199559P.
25-APR-2000; 2000WS-0199559P.
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03-MAY-2000; 2000US-0201516P.
17-MAY-2000; 2000US-0201516P.
22-MAY-2000; 2000WO-US013705.
32-MAY-2000; 2000WO-US014042.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-MAR-2000; 2000WO-US005841.
03-MAR-2000; 2000US-0187202P.
06-MAR-2000; 2000US-0186968P.
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2000WO-US030952
2000WO-US032678
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                                                                                                    18-DEC-2001 (first entry)
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05-JUN-2000;
28-JUL-2000;
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08-NOV-2000;
01-DEC-2000;
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98US-0084366P.
98US-0084414P.
98US-0084639P.
98US-0084640P.
98US-0084643P.
97US-0063541P.
97US-0063544P.
97US-0063564P.
97US-0063734P.
97US-0063870P.
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97US-0065311P.
97US-0066120P.
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970S-0066772P.
970S-0069335P.
970S-0069425P.
970S-0069870P.
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98US-0078939P.
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98US-0079786P.
98US-0080107P.
98US-0080194P.
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98US-0082568P.
98US-0082569P.
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98US-0080333P,
98US-0081049P.
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98US-0083495P.
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98US-0085582P.
98US-0085700P.
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98US-0088326P.
98US-0088167P.
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98US-0088212P.
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98US-0088655P.
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98US-0088826P.
98US-0088861P.
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98US-0086392P.
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98US-0077632P
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98US-0081195P
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98US-0082797P
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98US-0087098P
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98US-0087827P
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98US-0088824P
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27-MAR-1998;
31-MAR-1998;
31-MAR-1998;
28-OCT-1997;
28-OCT-1997;
29-OCT-1997;
31-OCT-1997;
31-OCT-1997;
21-NOV-1997;
24-NOV-1997;
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11-DEC-1997;
12-DEC-1997;
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18-DEC-1997;
10-MAR-1998;
11-MAR-1998;
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20-MAR-1998;
20-MAR-1998;
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01-APR-1998;
08-APR-1998;
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29-APR-1998;
05-MAY-1998;
06-MAY-1998;
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15-MAY-1998;
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29-APR-1998;
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15-MAY-1998;
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22-MAY-1998;
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                                                                                                                                                                                                                                                                          Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.

The PRO polypeptides and their associated nucleic acids can be used to detect the presence of a tumour in a mammal by comparing the level of expression of a PRO polypeptide in a test sample of cells from the animal and a control sample of normal cells, whereby a higher level of expression in the test sample indicates the presence of a tumour in the amimal. Mammals include dogs, cats, catle, horses, sheep, pigs, goats and rabbits but are preferably human. The polypeptides can be used to stimulate tumour necrosis factor (TMF) alpha release from human blood, when contacted with it. A specific polypeptide can be used to stimulate the proliferation or differentiation of chondrocyte cells. The PRO proteins can be used to determine the presence of tumours and also proteins can be used to determine the presence of tumours and also breast, prostate, rectal, cervical, or liver tumours, in mammalian subjects. The oligonucleotide probes specific for the PRO nucleic acids can be used for genetic analysis of individuals with genetic disorders
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                                                                                                                                                                       Novel nucleic acids encoding PRO polypeptides, used to diagnose the presence of tumors, such as prostate and breast tumors, in mammals and to screen for modulators of the compounds.
                                                                     Godowski PJ, Gurney AL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, PRO, cytostatic, tumour, cancer; breast, lung, stomach; liver, dog, cat; cow, horse, sheep, pig, goat, rabbit, ADEPT, antibody-dependent enzyme mediated prodrug therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30.0%; Score 6; DB 4; Length 1089; 100.0%; Pred. No. Se+02; ive 0; Mismatches 0; Indels
                                                            Goddard A, Godov
                                                                                   Wood WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU58532 standard; protein; 1089 AA
                                                                                                                                                                                                                                              Claim 11; Fig 266; 774pp; English.
                                                                 Chen J, Desnoyers L, ith V, Watanabe CK, W
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97US-0062250P.
97US-0063486P.
97US-0063120P.
97US-0063540P.
 20-DEC-2000; 2000WO-US034956.
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                                (GETH ) GENENTECH INC.
                                                                                                                      WPI; 2001-602746/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        687 WLRRYG 692
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 WLRRYG 17
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                                                                                     Smith V,
                                                                                                                                      N-PSDB; AAS46057
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24-0CT-1997;
24-0CT-1997;
28-0CT-1997;
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17-OCT-1997;
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                                                                   Baker KP,
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                                                                                       Pan J,
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PR 11.-UW-1998 98US-008865 PPR 11.-UW-1998 98US-008891 PPR 12.-UW-1998 98US-008991 PPR 17.-UW-1998 98US-008991 PPR 17.-UW-1998 98US-008991 PPR 17.-UW-1998 98US-008991 PPR 17.-UW-1998 98US-008991 PPR 18.-UW-1998 98US-008991 PPR 22.-UW-1998 98US-00990 PPR 22.-U
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Gaps
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Pred. No. 5e+02;
0; Mismatches 0
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llarity 100.0%; Pr
Conservative 0;
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97US-0059266P.
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97US-0063121P.
97US-0063540P.
97US-0063544P.
97US-0063544P.
97US-0063544P.
97US-0063544P.
97US-0063544P.
                     98US-0100930P.
98US-010104P.
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  98US-0100919P
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98US-00168978
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Matches 6; Conserv
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17-SEP-1998;
17-SEP-1998;
18-SEP-1998;
18-SEP-1998;
18-SEP-1998;
23-SEP-1998;
23-SEP-1998;
23-SEP-1998;
24-SEP-1998;
25-SEP-1998;
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30-SEP-1998
30-SEP-1998
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17-OCT-1997;
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31-OCT-1997;
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98US-0089538P
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27-MAY-1998

28-MAY-1998

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12. MAR-1998;

20. MAR-1998;

27. MAR-1998;

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31. MAR-1998;

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28-APR-1998;
29-APR-1998;
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21-APR-1998;
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22-APR-1998;
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29-APR-1998;
o'
                                                                                                                                                                                                                                                                                                                                                                                               Human; secreted and transmembrane protein; PRO; TNF-alpha; tumour necrosis factor alpha; chondrocyte cell; tumour; gene therapy; tissue typing.
                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                               0
                                                                                                                                                                                                         Similarity 100.0%; Score 6; DB 6; Length 1089; Similarity 100.0%; Pred. No. 5e+02; 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein (PRO) #133.
                                                                                                                                                                                                                                                                                                                     ABU84395 standard; protein; 1089 AA.
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970S-0053260P.
970S-0063212P.
970S-0063121P.
970S-0063121P.
970S-0063541P.
970S-0063541P.
970S-0063541P.
970S-0063734P.
970S-0063734P.
970S-0066120P.
970S-0069335P.
970S-0069335P.
        9805-0101472P

9805-0101475P

9805-0101738P

9805-0101739P

9805-0101743P

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9805-010224P

9805-010231P

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9805-010254P
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98US-0101471P
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Best Local Similarity
Matches 6; Conserv
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29-0CT-1997;
31-0CT-1997;
31-0CT-1997;
13-NOV-1997;
        23-SEP-1998;
23-SEP-1998;
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18-DEC-1997;
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98US-00904435P

98US-0090444P

98US-0090444P

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98US-0090540P

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10-SEP-1998;
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16-SEP-1998;
16-SEP-1998;
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                                                                                                                                                                                                                                                                                                  30.0%; Score 6; DB 6; I
100.0%; Pred. No. 5e+02;
.ive 0; Mismatches 0
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98US-0102240P.
98US-0102240P.
98US-0102331P.
98US-0102487P.
98US-0102571P.
98US-0102571P.
98US-0102684P.
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97US-0053266P.
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29-SEP-1998,
29-SEP-1998,
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30-SEP-1998,
10-OCT-1998,
02-OCT-1998,
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17 - OCT - 1997;

24 - OCT - 1997;

28 - OCT - 1997;

29 - OCT - 1997;

31 - OCT - 1998;

31 - OCT - 1
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                                                                                                           October 26, 2004, 07:22:37; Search time 65 Seconds (without alignments) 99.618 Million cell updates/sec
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1: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUBCOMB.pep:*} \)

2: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUBCOMB.pep:*} \)

3: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSC NEW PUB. pep:*} \)

4: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSC BWB PUB. pep:*} \)

5: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSC BWB PUB. pep:*} \)

6: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSC BWB PUB. pep:*} \)

7: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSC BWB PUB. pep:*} \)

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11: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSOB_R PUB. pep:*} \)

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15: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSOB_R PUB. pep:*} \)

16: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSD PUB. pep:*} \)

17: \( \text{cgn2} \) \( \text{Fptodata} \) 2 \( \text{Pubpaa} \) \( \text{VUSD PUB. pep:*} \)

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              GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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                                                                             - protein search, using sw model
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Gapop 60.0 , Gapext 60.0
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Jessen, Timm H.
Xu, Chanxing Wilson
TITLE OF INVENTION: INTERACTION TRAP SYSTEMS FOR DETECTING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLUM LIFE: FLODEY CLEAR
COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/162,538
FILING DATE: 04-Jun-2002
CLASSIFCATION: cUnknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/630,052
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: 08/504,538
FILING DATE: July 20, 1995
APPLICATION NUMBER: 08/278,082
FILING DATE: July 20, 1994
ATTORNEY/AGENT INFORMATION:
    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDBDNESS: NO. US20030113749A1 Relevant
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.0%; Score 13; DB 14; I
100.0%; Pred. No. 1.9e-06;
Live 0; Mismatches 0;
       Score 20; DB 14;
Pred. No. 1.4e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 00786/311001
                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Karen F. Lech
REGISTRATION NUMBER: 35,238
                                                                                                                                                                                                                                                                                                      INTERACTIONS
                                                                                                                                      RESULT 3
US-10-162-538-12
; Sequence 12, Application US/10162538
; Publication No. US20030113749A1
; GENERAL INFORMATION:
. APPLICANT: Brent, Roger
. McCOY, John M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
100.0%; Sco
100.0%; Pre
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                                                                                            1 OVWSSWALGWRWIRRYGWGM 20
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                                                                                                                                                                                                                                                                                                                                                                                   CITY: Boston
STATE: Massachusetts
                                                                    1 QVWSSWALGWRWLRRYGWGM
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          Query Match
Best Local Similarity 100.
Matches 20; Conservative
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Matches 13; Conserv
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| Sequence 2. Application No. US2030143626A1 |
| GENERAL INFORMATION: |
| APPLICANT: COLAS, PIERRE |
| APPLICANT: COLEN, PROGER |
| TILLE OF INVENTION: TARGETED MODIFICATION OF INTRACELLULAR COMPOUNDS |
| FILE REFERENCE: EGFP 3.0-015 |
| FILE REFERENCE: EGFP 3.0-015 |
| CURRENT APPLICATION UNMER: US/10/066,965A |
| CURRENT FILING DATE: 2002-12-09 |
| SOFTWARE: PATENTING OF 2.1 |
| SOFTWARE: PATENTING OF 2.1 |
| LENGTH: 20
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               Sequence
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, US-10-066-965A-8
                                                                                                                                  Sequence
                                                                                                                                                                                             Sequence
                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic ; OTHER INFORMATION: peptide US-10-066-965A-2
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            US-10-174-576-266
US-10-174-585-266
US-10-175-747-266
US-10-175-747-266
US-10-176-481-266
US-10-176-485-266
US-10-176-485-266
US-10-176-483-266
US-10-176-756-266
US-10-176-756-266
US-10-176-756-266
US-10-176-756-266
                                                                                                                                                               US-10-176-919-266
US-10-176-925-266
                                                                                                                                                                                                                                        ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 OVWSSWALGWRWLRRYGWGM 20
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Best Local Similarity 100.0%;
Matches 20; Conservative 0.
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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APPLICANT: Prendergast, John
APPLICANT: Goldstein, Neil I.

TITLE OP INVENTION: TARGET SPECIFIC SCREENING AND ITS USE FOR IDENTIFYING TARGET BINDE
FILE REPERRNCE: 2598-4009US1
CURRENT FILING DATE: 2002-10-24
PRIOR PILING DATE: 2001-10-24
NUMBER OF SEQ ID NOS: 537
SOFTWARE: Patentin version 3.1
SEQ ID NO 115
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Li, Ping,
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Clone ID: PAT_MRT4530_62377C.1.pep
US-10-437-963-163374
                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30.0%; Score 6; DB 16
100.0%; Pred. No. 37;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
30.0%; Score 6; DB 1
Best Local Similarity 100.0%; Pred. No. 23;
Matches 6; Conservative 0; Mismatches
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; Publication No. US20040123343A1
; GENERAL INFORMATION:
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Publication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: MISC_FEATURE
OTHER INFORMATĪON: DGI-2-20R-3-D10
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                  ORGANISM: Escherichia coli
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Best Local Similarity 100.
Matches 6; Conservative
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Publication No. US20030143626A1
FUBLICANT: USCOLAS, PIERRE
APPLICANT: COLAS, PIERRE
AUGUSTICANT COMPOUNDS
CURRENT FILING DATE: 202-12-09
NUMBER OF SEQ ID NOS: 33
SOCTWARE PATENTIN Ver. 2.1
SEQ ID NO 7
LENGTH: 20
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                 Sequence 1, Application US/10066965A
Publication No. US20030143626A1
GENERAL INFORMATION:
APPLICANT: COLAS, PIERRE
APPLICANT: COLAS, PIERRE
APPLICANT: COLES, PIERRE
APPLICANT: COLES, PIERRE
APPLICANT: COHEN, BARAK A.
TILLE OF INVENTION: TARGETED MODIFICATION OF INTRACELLULAR COMPOUNDS
FILE REFERENCE: EGYP 3.0-15
CURRENT APPLICATION NUMBER: US/10/066,965A
CURRENT FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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Pred. No. 1.9e-06;
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US-10-280-066-115
Sequence 115, Application US/10280066
Publication No. US20030180718A1
GENERAL INFORMATION:
APPLICANT: Pilutla, Renuka C.
APPLICANT: Spruyt, Michael
APPLICANT: Dedova, Olga
APPLICANT: Blume, Arthur J.
                                                                                                                                                                                                                                                                                                     SEQ ID NO 1
LENGTH: 20
TYPE: PRT
ORGANISM: Artificial Sequence
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Matches 13; Conserv
US-10-066-965A-1
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US-10-066-965A-7
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Gaps

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APPLICANT: Hyseq, Inc.
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: No. US20040053245Alel Nucleic Acids and Polypeptides
FILE REFERENCE: 21272-030
CURRENY APPLICATION NUMBER: US/10/276,774
CURRENY PILING DATE: 2000-11-18
PRIOR APPLICATION NUMBER: 09/560,875
PRIOR APPLICATION NUMBER: 09/560,875
PRIOR APPLICATION NUMBER: 09/496,914
PRIOR PILING DATE: 2000-04-27
PRIOR PILING DATE: 2000-06-27
PRIOR APPLICATION NUMBER: 09/496,914
PRIOR FILING DATE: 2000-02-03
SOFTWARE: Custom
SEQ ID NOS: 2700
SOFTWARE: LAST
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100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                            Length 112;
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APPLICANT: AKIYAM, YUTAKA
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: UP 2001/246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
SOFTWARE: PATENTIN VET. 2.1
                                                                                                                                                                                                                                                            DB 9;
                                                                                                                                                                                                                                                            30.0%; Score 6; DB 9, 100.0%; Pred. No. 86; rative 0; Mismatches
CURRENT FILING DATE: 2001-09-20
PRIOR APPLICATION NUMBER: USSN 60/208,427
PRIOR FILING DATE: 2000-05-30
NUMBER OF SEQ ID NOS: 2125
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1614
LENGTH: 112
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Sequence 1740, Application US/10017161
; Publication No. US20030143668A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 1454, Application US/10276774; Publication No. US20040053245A1; GENERAL INFORMATION:
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Best Local Similarity 100...
6; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens
US-10-276-774-1454
                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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US-10-276-774-1454
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Patent No. US20020082206A1

GENERAL INFORMATION:
APPLICANT: Leach, Martin D.
APPLICANT: Conley, Pamela
APPLICANT: Taw, Debbie
APPLICANT: Topper, James
APPLICANT: Thereby
TITLE OF INVENTION: Thereby
FILE REPERENCE: 21402-013 (Gura-313)
CURRENT APPLICATION NUMBER: US/09/867,550
                                                                           APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPREBUCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 136125
LENGTH: 65
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Sequence 252446, Application US/10424599

Publication No. US20040031072A1

GENERAL INFORMATION:

APPLICANT: La Rosa Thomas J

APPLICANT: Zhou Yihua

APPLICANT: Cao Yongwei

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(33223)B

CURRENT APPLICATION UNMERR: US/10/424,599

CURRENT FILING DAIE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684
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100.0%; Pred. No. 64;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                       30.0%; Score 6; DB 16; Length 65; 100.0%; Pred. No. 58; 0; Indels tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Clone ID: PAT_MRT4530_37734C.1.pep US-10-437-963-136125
       Wu, Wei
Boukharov, Andrey A.
Barbazuk, Brad
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Matches 6; Conservative
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ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Glycine max
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Best Local Similarity
Matches 6; Conserv
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LENGTH: 74
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US-09-867-550-1614
         APPLICANT:
APPLICANT:
APPLICANT:
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US-10-017-161-1740

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APPLICANT: La Novalitation:
APPLICANT: La Novalic, David K.
APPLICANT: Stool, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Li, Ping
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE REFERENCE: 39-21(5321)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
LENGTH: 202
                                                                                                                                                  Sequence 40080, Application US/10425114
| Publication No. US2004003488A1
| Publication No. US2004003488A1
| GENERAL INFORMATION:
| APPLICANT: Liu, Jingdong
| APPLICANT: Zhou, Yihua
| APPLICANT: Zhou, Yihua
| APPLICANT: Zhou, Yihua
| APPLICANT: Tabaska, Jack E
| APPLICANT: Tabaska, Jack E
| APPLICANT: Cao, Yongwei
| TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
| TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
| TITLE REFERENCE: 38-21(53313)B
| CURRENT APPLICATION NUMBER: US/10/425,114
| CURRENT FILING DATE: 2003-04-28
| NUMBER OF SEQ ID NOS: 73128
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_9481C.1.pep
US-10-437-963-199236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 15; Le:
o. 1.3e+02;
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LOCATION: (1)..(202)
OTHER INFORMATION: unsure at all Xaa locations
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100.0%; Pred. No. 1.3
tive 0; Mismatches
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Publication No. US20040123343A1
GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 6; Conservative
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ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
      LRRYGW 18
                                             72 LRRYGW 77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 16
US-10-437-963-199236
                                                                                                                                      US-10-425-114-40080
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LENGTH: 196
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US-10-425-114-43111

Sequence 43111, Application US/10425114

Publication No. US20040034888A1

GENERAL INFORMATION:

APPLICANT: Liu, Jingdong

APPLICANT: Zhou, Yihua

APPLICANT: Acreen, Screen, Steven E

APPLICANT: APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

CURRENT APPLICANTON NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 73128

LENGTH: 180
                                                                                                                                                                                                                                      Sequence 1396, Application US/10292798

Sequence 1396, Application US/10292798

Publication No. US2003235833A1

GENERAL INFORMATION:

APPLICANT: SUWA, MAXIXO

APPLICANT: ASAI, KIVOSHI

APPLICANT: ASAI, KIVOSHI

TITLE OF INVENTION: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS

FILE REFERENCE: 084335/166

CURRENT PALLOAN NUMBER: US/10/292,798

CURRENT FILING DATE: 2002-11-13

PRIOR FILING DATE: 2001-12-18

PRIOR FILING DATE: 2001-06-18

NUMBER OF SEQ ID NOS: 2070

SOFTWARE: PatentIN Ver. 2.1

SEQ ID NO 1396

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                         Length 165;
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                                                               0; Indels
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                  DB 14; Le
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les 0;
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US-10-425-114-43111
30.0%; Scor.
100.0%; Pred. No. ...
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100.0%;
Query Match
Best Local Similarity 100.
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Best Local Similarity 100.v
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Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
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ORGANISM: Glycine max
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                                                                                                                                                                                                           RESULT 13
US-10-292-798-1396
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US-10-383-317-16
| Sequence 16, Application US/10383317
| Publication No. US20040001849A1
| Publication No. US20040001849A1
| Publication No. US20040001849A1
| APPLICANT: Punnonen, Unha
| APPLICANT: Bass Steven H. |
| APPLICANT: Whalen, Robert Gerald APPLICANT: Howard, Russell N. |
| PILE REFERENCE: 01809-02810|
| PRIOR PILING DATE: 1999-02-10|
| PRIOR PLING DATE: 1999-02-10|
| PRIOR FILING DATE: BARLIER FILING DATE: 1998-02-11|
| PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-11|
| PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-11|
| PRIOR PRIOR APPLICATION NUMBER: EARLIER FILING DATE: 1998-10-23|
| NUMBER OF SEQ ID NOS: 25
| SOFTWARE: PatentIn Ver. 2.0|
| SEQ ID NO 16|
| INNOVINGER NOSE 16|
| I
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                                                                                   0;
                       Length 243;
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                    30.0%; Score 6; DB 16; Length 243
100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Punnonen, Juha
APPLICANT: Bass, Steven H.
APPLICANT: Bass, Steven H.
APPLICANT: Whalen, Robert Gerald
APPLICANT: Whalen, Robert Gerald
APPLICANT: Howard, Russell
APPLICANT: Howard, Willem P.C.
APPLICANT: Stemmer, Willem P.C.
TITLE OF INVENTION: Antigen Library Immunization
TITLE OF INVENTION: Antigen Library Immunization
CURRENT APPLICATION NUMBER: US/09/247,890
CURRENT FILING DATE: 1999-02-10
EARLIER PPLICATION NUMBER: US 60/074,294
EARLIER FILING DATE: 1998-02-11
FEARLIER PILING DATE: 1998-02-13
NUMBER OF SEQ ID NOS: 25
SOFFWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30.0%; Score 6; DB 9; Len. 100.0%; Pred. No. 1.7e+02; tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Woodchuck hepatitis B virus
                                                                                                                                                                                                                                                                                                                                              US-09-247-890-16; Sequence 16, Application US/09247890; Publication No. US20020198162A1; GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conservat
210 SSWALG 215
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LENGTH: 282
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; Sequence 13424, Application US/10437963
; Publication No. US20040123343A1
; Deblication No. US20040123343A1
; Deblication No. US20040123343A1
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Cao, Vongwei
; APPLICANT: Cao, Yongwei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Bucknarov, Andrey A.
; APPLICANT: Brabazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; TITLE OF INVENTION NUMBER: US/10/437,963
; CURRENT APPLICATION NUMBER: US/10/437,963
; SEQ ID NO 13424
; LENGTH: 240
; LENGTH: 240
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APPLICANT: Li, Fing
APPLICANT: Li, Fing
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPREMENTE: 38-21(53221) B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 199246
LENGTH: 243
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30.0%; Score 6; DB 16; Length 240;
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
      Indels
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US-10-437-963-134324
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OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
          0; Mismatches
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Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
             6; Conservative
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ORGANISM: Oryza sativa
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                                                                                                                                     178 LGWRWL 183
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                                                                            8 LGWRWL 13
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                                                                                                                                                                                                                                      RESULT 17
US-10-437-963-134324
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             Matches
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TELECOMMUNICATION INFORMATION:
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ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2, Application US/09812862 Patent No. US20020035081A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGIH: 346 amino acids
      CURRENT FILING DATE: 2003-04-28
WUMBER OF SEQ ID NOS: 73128
SEQ ID NO 37645
LENGTH: 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                     Query Match
Best Local Similarity 100.0
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                                                                                                                                                                                                                                                                                                                                                       52 SSWALG 57
                                                                                                                                                                                                                                                                                                              4 SSWALG 9
                                                                                              TYPE: PRT
ORGANISM: Zea mays
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US-09-812-862-2
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APPLICANT: Tabaska, Jack B
APPLICANT: Tabaska, Jack B
APPLICANT: Cao, Yongwa B
APPLICANT: Cao, Yongwa C
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
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NAME/KEY: SITE
LOCATION: (29)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: SITE
LOCATION: (157)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
LOCATION: (205)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
OTHER INFORMATION: As equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: SITE
LOCATION: (275)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (290)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids US-09-764-875-621
                                                                                         Gaps
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                                                                                                                                                                                                                                                                        Sequence 621, Application US/09764875;
Publication No. US20040018969A1;
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: D4202
CURRENT APPLICATION NUMBER: US/09/764,875;
CURRENT PILING DATE: 2001-01-17;
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1249;
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: PATENTING D515;
LENGTH: 305
TYPE: PRT
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                                                 Length 282;
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                                                                                         0; Indels
                                            DB 15; Le
                                        30.0%; Score 6; DB 1
100.0%; Pred. No. 1.7
tive 0; Mismatches
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Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Rovalic, David K.
APPLICANT: Rovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
                    Query Match
Best Local Similarity luv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                   210 SSWALG 215
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                               4 SSWALG 9
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                                                                                                                                                                                                                                    RESULT 21
US-09-764-875-621
US-10-383-317-16
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Gaps
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                                                                                                                                                                                                                                         Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                         Length 324;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
30.0%; Score 6; DB 9; Length 346;
Best Local Similarity 100.0%; Pred. No. 2e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
                                                                       Indels
                                            DB 15; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/282001
; OTHER INFORMATION: Clone ID: 700264190_FLI.pep
US-10-425-114-37645
                                           30.0%; Score 6; DB 1
100.0%; Pred. No. 1.9
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                         ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
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Sequence 4601, Application US/10369493

Publication No. US20030233675A1

GENERAL INFORMATION:

APPLICANT: Cao, Yongwei

APPLICANT: Gladen, Barry S.

APPLICANT: Gladen, Barry S.

APPLICANT: Chen, Kianfeng

TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROFERIES

TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES

TITLE OF LING DATE: 18-10 (52052) B

CURRENT APPLICATION NUMBER: US/10/369,493

CURRENT APPLICATION NUMBER: US/10/369,493

FRICK APPLICATION NUMBER: US/10/369,493

PRIOR FILING DATE: 2002-02-21
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                                                                                                                                                                                                                                                         APPLICANT: Tarwick, Journ
APPLICANT: Yamamoto, Robert
APPLICANT: Xu, H.
TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REFRENCE: ELITRA.034A
CURRENT APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR PLILNG DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-09-06
PRIOR APPLICATION NUMBER: 60/230,347
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-110-23
PRIOR FILING DATE: 2000-110-23
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR PRILING DATE: 2000-12-20
PRIOR FILING DATE: 2000-12-20
PRIOR PRILING DATE: 2000-12-20
PRIOR PRILING DATE: 2000-12-20
PRIOR PRILING DATE: 2001-02-09
PRIOR FILING DATE: 2001-02-09
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Publication No. US20040029129A1
                                                   APPLICANT: Wang, Liangsu
APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
Haselbeck, Robert
APPLICANT: Ohlsen, Kari
                                                                                                                                                                                          Zyskind, Judith
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Trawick, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Sequence 160000, Application US/10424599

Publication No. US20040031072A1

GENERAL INFORMATION:

APPLICANT: La Rosa Thomas J

APPLICANT: Con Yongwei

APPLICANT: Con Yongwei

APPLICANT: Cao Yongwei

TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REPRESENCE: 38-1(53223)

CURRENT APLICATION NUMBER: US/10/424,599

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 16000

LENGTH: 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30.0%; Score 6; DB 15; Length 369; 100.0%; Pred. No. 2.1e+02;
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                                                                                                                                 Sequence 4, Application US/10343650A
; Sequence 4, Application No. US20040067499A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TILLE OF INVENTION: NOVEL G-PROTEIN COUPLED RECEPTOR
GENERAL PAPPLICATION NUMBER: US/10/343,650A
CURRENT APPLICATION NUMBER: US/200/237818
; PRIOR APPLICATION NUMBER: UP 2000/237818
; PRIOR APPLICATION NUMBER: UP 2001/34434
; PRIOR APPLICATION NUMBER: UP 2001/34434
; PRIOR FILING DATE: 2000-08-04
; PRIOR APPLICATION NUMBER: UP 2001/34434
; NUMBER OF SEQ ID NOS: 694
; SOFTWARE: PatentIn Ver. 2.1
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US-10-424-599-160000
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OTHER INFORMATION: unsure at all Xaa locations
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Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity
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        274 SSWALG 279
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US-10-282-122A-47856
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US-10-424-599-160000
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                                                                                           RESULT 24
US-10-343-650A-4
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LENGTH: 368
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-190-115-111
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; ORGANISM: Homo sapiens
US-10-190-115-114
                                                                                       Query Match
Best Local Similarity
Local 6; Conserv?
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SEQ ID NO 114
LENGTH: 401
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/190,115
CURRENT FILING DATE: 2003-02-10
PRIOR PLING DATE: 2003-02-04
PRIOR PLING DATE: 2002-04-01
PRIOR PLING DATE: 2002-04-01
PRIOR APPLICATION NUMBER: 60/386,816
PRIOR PLING DATE: 2002-04-07
PRIOR PLING DATE: 2000-07-03
PRIOR PLING DATE: 2000-07-07
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SOFTWARE: CuraSeqList version 0.1
SEQ ID NO 111
LENGTH: 401
                                                                                                                                                                                                                                       .
0
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                                                                                                                                                                                                                                  0; Indels
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Best Local Similarity 100.0%; Pred. No. 2.2
Matches 6; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 111, Application US/10190115
Publication No. US20030207394A1
GENERAL INFORMATION:
APPLICANT: Boldog, Ferenc L.
APPLICANT: Burgess, Catherine E.
APPLICANT: Gasman, Stacke J.
APPLICANT: Grosse, William M.
APPLICANT: Grosse, William M.
APPLICANT: Gusev, Vladimir Y.
APPLICANT: Lepley, Denise M.
APPLICANT: Lepley, Denise M.
APPLICANT: Liepley, Denise M.
APPLICANT: Liepley, Amanda J.
APPLICANT: Mezick, Amanda J.
APPLICANT: Padigaru, Muralidhara
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Shimkets, Richard A.
Spaderna, Steven K.
Spytek, Kimberly A.
Szekeres, Edward S. Jr.
Taupier, Raymond J. Jr.
Tcherney, Velizar T.
                                                                          TYPE: PRT
ORGANISM: Burkholderia fungorum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patturajan, Meera
Rastelli, Luca
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 4601
LENGTH: 400
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Gaps

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### PRIOR PRINCENTION NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME FILE REPERENCE: 21402-050 CIP (CURRENT APPLICATION NUMBER: US/10/190,115 CURRENT APPLICATION NUMBER: US/10/190,115 CURRENT FILING DATE: 2003-02-10 PRIOR PAPLICATION NUMBER: 60/303,168 PRIOR FILING DATE: 2002-04-01 PRIOR PELING DATE: 2002-04-01 PRIOR APPLICATION NUMBER: 60/366,816 PRIOR FILING DATE: 2002-06-07 PRIOR PELING DATE: 2000-07-03 PRIOR PAPLICATION NUMBER: 60/215,854 PRIOR PELING DATE: 2000-07-03 PRIOR PELING DATE: 2000-07-07 PRIOR PEL
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30.0%; Score 6; DB 14; Length 401;
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
Indels
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    Mismatches
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Publication No. US20030207394A1
GENERAL INFORMATION:
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APPLICANT: Boldog, Ferenc L.
APPLICANT: Burgess, Catherine B.
APPLICANT: Casman, Stacie J.
APPLICANT: Grosse, William M.
APPLICANT: Gusev, Vladimir Y.
APPLICANT: Ji, Weizhen
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Szekeres, Edward S. Jr.
Taupier, Raymond J. Jr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Padigaru, Muralidhara
Patturajan, Meera
Rastelli, Luca
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Zerhusen, Bryan D.
Voss, Edward Z.
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Liu, Xiachong
Mezick, Amanda J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Suresh G.
    6; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                   60 WSSWAL 65
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                                                                                 3 WSSWAL 8
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APPLICANT:
        Matches
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                                                                                                                                         Gaps
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 136
SOFTWARE: CuraSeqList version 0.1
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                                               30.0%; Score 6; DB 14; Length 401; 100.0%; Pred. No. 2.2e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 116. Application US/10190115
Publication No. US20030207394A1
GENERAL INFORMATION:
APPLICANT: Alsobrook, John P. II
APPLICANT: Boldog, Ferenc L.
APPLICANT: Boldog, Ferenc L.
APPLICANT: Gasse, William M.
APPLICANT: Grosse, William M.
APPLICANT: Grosse, William M.
APPLICANT: Grosse, William M.
APPLICANT: Lepley, Denise M.
APPLICANT: Padigaru, Muralidhara, APPLICANT: Padigaru, Muralidhara, APPLICANT: Patturajan, Meera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Spytek, Kimberly A.
Szekeres, Edward S. Jr.
Taupier, Raymond J. Jr.
Tchernev, Velizar T.
Zerhusen, Bryan D.
Voss, Edward Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shenoy, Suresh G.
Shimkets, Richard A.
Spaderna, Steven K.
                                                                                                                                         6; Conservative
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ORGANISM: Homo sapiens
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Best Local Similarity
                                                            Query Match
Best Local Similarity
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APPLICANT:
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369 LRRYGW 374

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WSSWAL 65

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APPLICANT: Li, Ping,
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: 1032131
TITLE REFRENCE: 38-21321)8
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 122935
LENGTH: 500
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APPLICANT: Turano, Frank J.
TITLE OF INVENTION: Methods for Regulating Plant GABA Production FILE REPERENCE: 7224-62
CURRENT PILING DATE: 2002-07-01
PRIOR APPLICATION NUMBER: US 60/246,367
PRIOR FILING DATE: 2000-11-07
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.1
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30.0%; Score 6; DB 16; Length 500;
Best Local Similarity 100.0%; Pred. No. 2.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
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30.0%; Score 6; DB 14; Length 502
Best Local Similarity 100.0%; Pred. No. 2.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Clone ID: PAT_MRT4530_25818C.l.pep US-10-437-963-122935
                                                          Sequence 122935, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION.
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US-10-289-161A-9
; Sequence 9, Application US/10289161A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-006-852-2; Sequence 2, Application US/10006852; Publication No. US20030046732A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                               Boukharov, Andrey A.
Barbazuk, Brad
                                                                                                                          APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-10-006-852-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          397 LRRYGW 402
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                                    US-10-437-963-122935
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LENGTH: 502
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Fublication No. US20040123343A1

GENERAL INPORMATION:
GENERAL INPORMATION:
APPLICANT: AROSa, Thomas J.
APPLICANT: Cao, Yonowei
APPLICANT: Cao, Yonowei
APPLICANT: Barbazuk, Brad
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICANTON NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14

NUMBER OF SEQ ID NOS: 204966

SEQ ID NO 122938
                          Sequence 7358, Application US/10369493

Sequence 7358, Application US/10369493

Publication No. US20030233675A1

GENERAL INFORMATION:

APPLICANT: Cao, Yongwei

APPLICANT: Goldman, Barry S.

APPLICANT: Goldman, Barry S.

APPLICANT: Goldman, Barry S.

TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF FILE REFERENCE: 38-10 (52052)B

CURRENT APPLICATION NUMBER: US/10/369,493

CURRENT APPLICATION NUMBER: US/10/369,493

CURRENT APPLICATION NUMBER: US 60/360,039

PRIOR FILING DATE: 2002-02-21

NUMBER OF SEQ ID NOS: 47374

SEQ ID NO 7358
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US-10-437-963-122938
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CTHER INFORMATION: unsure at all Xaa locations US-10-369-493-7358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Burkholderia cepacia FEATURE:
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Best Local Similarity 100.
Matches 6, Conservative
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ORGANISM: Oryza sativa
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Best Local Similarity
Matches 6; Conserv
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RESULT 32
US-10-369-493-7358
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235 LGWRWL 240
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Matches 6; Conserv
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Best Local Similarity
Matches 6; Conserv
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                                                                   ; ORGANISM: Rai
US-10-308-163-16
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SEQ ID NO 16
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                                                                 APPLICANT: Silos-Santiago, Inmacculada
TITLE OF INVENTION: Methods and Compositions to Treat Pain
TITLE OF INVENTION: and Painful Disorders Using 577, 20739 or 57145
FILE DEPINVENTION: and Painful Disorders Using 577, 20739 or 57145
FILE OF INVENTION: and Painful Disorders Using 577, 20739 or 57145
FILE OF INVENTION: and Painful Disorders Using 577, 20739 or 57145
CURRENT APPLICATION NUMBER: US/10/289,161A
PRIOR FILING DATE: 2001-11-06
NUMBER: OF SEQ ID NOS: 9
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                              30.0%; Score 6; DB 14; Length 548; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/10308163
; Sequence 8, Application US/10308163
; Publication No. US20040106147A1
; GENERAL INFORMATION:
; APPLICANT: UNCKA, Karl
; APPLICANT: UNCKA, Karl
; APPLICANT: FUKS, Baruno
; TITLE OF INVENITON: Levetiracetam Binding Site
; TITLE OF INVENITON: Levetiracetam Binding Site
; TILE REFERENCE: 53529-5007
; CURRENT APPLICATION NUMBER: US/10/308,163
; CURRENT APPLICATION NUMBER: US/10/308,163
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: PatentIN version 3.1
; SEQ ID NO 8
INNEGHT: 548
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; Publication No. US20040106147A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: UNCH, Berkley
; APPLICANT: UNCKA, Karl
APPLICANT: WOCKA, Karl
APPLICANT: EUKS, Bruno
; TITLE REFERENCE: 53529-5007
; CURRENT PILING NUMBER: US/10/308,163
; CURRENT FILING DATE: 2002-12-03
; NUMBER OF SEQ ID NOS: 16
                           GENERAL INFORMATION:
APPLICANT: Millennium Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin version 3.1
         US20030152970A1
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Best Local Similarity 100.
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US-10-308-163-8
                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo Sapiens
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Best Local Similarity
Matches 6; Conserv
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US-10-308-163-16
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                                                                                                                                                                                                                                                                                                                LENGTH: 548
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US-10-308-163-8
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US=1U-/25-1By-8

Sequence 8, Application US/10725189

Sequence 8, Application US/2040204388A1

Sequence 8, Application No. US20040204388A1

GENERAL INFORMATION:

APPLICANT: UEVEN. S.A.

APPLICANT: LYNCH, Berkley

APPLICANT: LYNCH, Berkley

APPLICANT: NOCKA, Karl

APPLICANT: NOCKA, Karl

APPLICANT: RINCH Seizures, neurological diseases, endocrinopathies and hormonal;

TITLE OF INVENTION: Seizures, neurological diseases, endocrinopathies and hormonal;

TITLE OF INVENTION: Seizures, 120,10,725,189

CURRENT APPLICATION NUMBER: US/10/725,189

FILE REFERENCE: 53529-5007-01-05

PRIOR APPLICATION NUMBER: US 60/430,372

PRIOR APPLICATION NUMBER: US 60/430,372

PRIOR APPLICATION NUMBER: US 60/506,764

PRIOR FILING DATE: 2003-12-03

NUMBER OF SEQ ID NOS: 16

NUMBER OF SEQ ID NOS: 16
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APPLICANT: INNCH, Berkley
APPLICANT: INNCH, Berkley
APPLICANT: INNCH, Berkley
APPLICANT: MOCKA, Karl
APPLICANT: WOCKA, Warl
TITLE OF INVENTION: Methods for the identification of agents for the treatment of
TITLE OF INVENTION: Methods for the identification of adents and hormonal
TITLE OF INVENTION: 41seases
FILE REFERENCE: 53529-5007-01-US
CURRENT APPLICATION NUMBER: US 60/430,372
PRIOR FILING DATE: 2003-12-03
PRIOR FILING DATE: 2003-12-03
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1
SEQ ID NO 16
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                                                                                                       Length 548;
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Pred. No. 2.8e+02;
0; Mismatches 0;
                                                                                                       30.0%; Score 6; DB 16; Lt 100.0%; Pred. No. 2.8e+02;
                                                                                                                                                                0; Mismatches
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100.0%; Pre
TYPE: PRT ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                6; Conservative
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APPLICANT: La Rosa, Thomas J.
APPLICANT: La Rosa, Thomas J.
APPLICANT: La Lou, Yahua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Bunkharov, Andrey A.
APPLICANT: Bunkharov, Andrey A.
APPLICANT: Brabazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53.221) B
CURRENT APPLICATION NUMBER: US/10/437,963
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                                                                                                                                                                                                                                     LOCATION: (1)..(617)
OTHER INFORMATION: unsure at all Xaa locations
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OTHER INFORMATION: unsure at all Xaa locations
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    FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 117634
LENGTH: 617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 144980, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
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Patent No. US20020098491A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 144980
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Best Local Similarity 100.
Matches 6; Conservative
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APPLICANT: Wang, Xiaoming
APPLICANT: Scoville, John
                                                                                                                                               TYPE: PRT
ORGANISM: Oryza sativa
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ORGANISM: Oryza sativa
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Best Local Similarity
Matches 6; Conserv
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NAME/KEY: unsure
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US-10-768-158-24

US-10-768-158-24

Sequence 24, Application US/10768158

Fublication No. US20040204359A1

GENERAL INFORMATION:

APPLICANT: Milennium Pharmaceuticals, Inc.

APPLICANT: Karicheti, Venkateswarlu

APPLICANT: Eliasof, Scott D.

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING

TITLE OF INVENTION: PAIN AND PAINFUL DISORDERS USING 16386, 15402, 21165, 1423,

TITLE OF INVENTION: 94921, 16852, 33260, 58573, 30911, 85913, 14303, 16816,

TITLE OF INVENTION: 17827 OR 23620

TITLE OF DATE: 2004-01-20180NOMIM

CURRENT APPLICATION NUMBER: US/10/768158
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
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                                                                                                                30.0%; Score 6; DB 17; Length 548; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/444,781
PRIOR FILING DATE: 2003-02-04
PRIOR FILING DATE: 2003-02-04
PRIOR PILING DATE: 2003-05-05
PRIOR PILING DATE: 2003-03-05
PRIOR PLING DATE: 2003-03-13
PRIOR PLING DATE: 2003-03-13
PRIOR PILING DATE: 2003-04-16
PRIOR PILING DATE: 2003-04-16
PRIOR PILING DATE: 2003-06-16
PRIOR FILING DATE: 2003-06-16
PRIOR FILING DATE: 2003-07-30
NUMBER OF SEQ ID NOS: 46
SOFTWARE: FASTSEQ for Windows Version 4.0
SENGTH: 548
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Publication No. US2004012334341
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yihua
APPLICANT: Cao, Yihua
APPLICANT: Ru, Wei
APPLICANT: Buukharov, Andrey A.
APPLICANT: Barbaruk, Brad
; IENGTH: 548
; TYBE: PRT
; ORGANISM: Rattus norvegicus US-10-725-189-16
                                                                                                                                                                6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                     235 LGWRWL 240
                                                                                                                                   Best Local Similarity
Matches 6; Conserv
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Best Local Similarity
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GENERAL INFORMATION:

APPLICANT: Walke, D. Wade

APPLICANT: Walke, John

APPLICANT: Walke, John

APPLICANT: Turner, C. Alexander Jr.

APPLICANT: Turner, C. Alexander Jr.

APPLICANT: Turner, C. No. US20020098491A1el Human Semaphorin Homologs and Polynucleotid

FILE REFERENCE: LEX-0177-USA

CURRENT APPLICATION NUMBER: US 60/205,274

PRIOR PRING DATE: 2000-06-02

NUMBER OF SEQ ID NOS: 50

SOFTWARE: FastSEQ for Windows Version 4.0
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100.0%; Pred. No. 4.4
tive 0; Mismatches
PRIOR APPLICATION NUMBER: US 60/205,274
PRIOR FILING DATE: 2000-05-18
PRIOR FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 50
SOFTWARE: PASLSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 1034
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Best Local Similarity 100.
Matches 6; Conservative
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US-09-854-845-8
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                                                                                                                                                         TYPE: PRT
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Sequence 14, Application US/09854845

Patent No. US20020038491A1

GENERAL INFORMATION:
APPLICANT: Walk, D. Wade

APPLICANT: Turner, C. Alexander Jr.
APPLICANT: Turner, C. Alexander Jr.
TITLE OF INVENTION: NO. US20020098491A1e1 Human Semaphorin Homologs and Polynucleotid
FILE REFERENCE: LEX-0177-USA

CURRENT APPLICATION NUMBER: US/09/854,845

CURRENT APPLICATION NUMBER: US/09/854,845

CURRENT APPLICATION NUMBER: US/0205,274

PRIOR FILING DATE: 2000-05-18

PRIOR FILING DATE: 2000-06-05

NUMBER OF SEQ ID NOS: 50

NUMBER OF SEQ ID NOS: 50

SOFTWARE: FastSEQ for Windows Version 4.0
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TITLE OF INVENTION: No. US20020098491Alel Human Semaphorin Homologs and Polynucleotid
FILE REFERENCE: LEX-0177-USA
CURRENT PAPLICATION NUMBER: US/09/854,845
CURRENT PAPLICATION NUMBER: US 60/205,274
PRIOR APPLICATION NUMBER: US 60/205,274
PRIOR APPLICATION NUMBER: US 60/208,893
PRIOR APPLICATION NUMBER: US 60/208,893
PRIOR FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16
LENGTH: 939
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                                                                                                                                                                                                                                                                                                                             Length 939;
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100.0%; Pred. No. 4.1
Live 0; Mismatches
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Matches 6; Conservative
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ORGANISM: homo sapiens
                                                                                                                                                                                                                                                      TYPE: PRT
CORGANISM: homo sapiens
US-09-854-845-16
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Matches 6; Conserv
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LENGTH: 954
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PRIOR APPLICATION NUMBER: 60/093763
PRIOR PLING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/09912
PRIOR PLING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/09912
PRIOR PLING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/09915
PRIOR PLING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-15
PRIOR PLING DATE: 1998-09-15
PRIOR PLING DATE: 1998-09-16
PRIOR PLING DATE: 1998-09-17
PRIOR PLING DATE: 1998-09-18
PRIOR PLING DATE: 1998-09-18
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APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Stewart, Timothy A.
APPLICANT: Tunas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Williams, P. Mickey
APPLICANT: Williams, P. Mickey
APPLICANT: Williams, P. Mickey
APPLICANT: William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Secreted and Transmembrane
FILE REFERENCE: P2830PlC1
CURRENT APPLICATION NUMBER: 60/098716
PRIOR APPLICATION NUMBER: 60/09872
PRIOR APPLICATION NUMBER: 60/09872
PRIOR APPLICATION NUMBER: 60/09872
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/09875
PRIOR APPLICATION NUMBER: 60/09875
PRIOR FILING DATE: 1998-09-01
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PRIOR APPLICATION NUMBER: 60/098750
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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 8 LENGTH: 1078 TYPE: PRT
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Publication No. US20030073129A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Bostetan, David
APPLICANT: Desnoyers, Luc
APPLICANT: Besnoyers, Luc
APPLICANT: Desnoyers, Luc
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Grimaldi, Christopher J.
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APPLICATION NUMBER: 60/099741
FILING DATE: 1998-09-10
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FILING DATE: 1998-09-02
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FILING DATE: 1998-09-02
APPLICATION NUMBER: 60/098843
FILING DATE: 1998-09-02
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FILING DATE: 1998-09-09
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APPLICATION NUMBER: 60/099642
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Fong, Sherman
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Gao, Wei-Qiang
Goddard, Audrey
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                                                                                                         ; ORGANISM: homo sapiens
US-09-854-845-8
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Best Local Similarity
Matches 6, Conserv
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Length 1089;

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PRIOR PEDICATION NUMBER: 60/102107
PRIOR APPLICATION NUMBER: 60/102207
PRIOR PELING DATE: 1998-09-24
PRIOR PLING DATE: 1998-09-29
PRIOR PELING DATE: 1998-09-29
PRIOR PLING DATE: 1998-09-30
PRIOR PLING DATE: 1998-10-01
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PRIOR PLING DATE: 1998-10-01
PRIOR PLING DATE: 1998-10-07
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R APPLICATION NUMBER: 60/101916
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APPLICATION NUMBER: 60/105694
FILING DATE: 1998-10-26
APPLICATION NUMBER: 60/105807
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APPLICANT: SILLY VICTORIA A
APPLICANT: STATE, VICTORIA A
APPLICANT: MARIANDE, COLIN K
APPLICANT: WARGANDE, COLIN K
APPLICANT: WARGANDE, MILLIAM I.
APPLICANT: WARGANDE, MILLIAM I.
APPLICANT: WARGANDE, COLIN K
FILLE OF INVENTION: SECRETED AND TRANSMENRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: SECRETED AND TRANSMENRANE POLYPEPTIDES AND NUCLEIC
CURRENT PILLING DATE: 1949 MRZE: 40/65263
PRIOR PELLING DATE: 1970-018
PRIOR PELLING DATE: 1970-018
PRIOR PELLING DATE: 1970-018
PRIOR PELLING DATE: 1970-10-27
PRIOR PELLING DATE: 1970-10-28
PRIOR PELLING DATE: 1970-10-29
PRIOR PELLING DATE: 1970-10-29
PRIOR PELLING DATE: 1970-10-31
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30.0%; Score 6; DB 10; Lo
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ive 0; Mismatches 0;
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FILING DATE: 1997-12-18
APPLICATION NUMBER: 60/077450
              30.08; Pre
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Watanabe, Colin K.
Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
                                                              6; Conservative
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                                                                                                                                                                       687 WLRRYG 692
                          Best Local Similarity
Matches 6; Conserv
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US-10-052-586-266
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APPLICANT:
APPLICANT:
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APPLICANT:
           Query Match
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DR FILING DATE: 1998-04-09

DR APPLICATION NUMBER: 60/081838

R FILING DATE: 1998-04-15

R APPLICATION NUMBER: 60/082568

R FILING DATE: 1998-04-21

R APPLICATION NUMBER: 60/082569

R FILING DATE: 1998-04-21
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PRIOR FILING DATE: 1938-05-07
PRIOR PRING DATE: 1938-05-07
PRIOR PELING DATE: 1998-05-15
PRIOR PELING DATE: 1998-05-15
PRIOR APPLICATION NUMBER: 60/085579
PRIOR APPLICATION NUMBER: 60/08558
PRIOR APPLICATION NUMBER: 60/08558
PRIOR PILING DATE: 1998-05-15
PRIOR PAPLICATION NUMBER: 60/08580
PRIOR PELING DATE: 1998-05-15
PRIOR APPLICATION NUMBER: 60/085700
PRIOR APPLICATION NUMBER: 60/085700
PRIOR APPLICATION NUMBER: 60/085700
PRIOR PELING DATE: 1998-05-15
PRIOR PELING DATE: 1998-05-15
PRIOR PELING DATE: 1998-05-15
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R APPLICATION NUMBER: 60/083496
R FILING DATE: 1998-04-29
R APPLICATION NUMBER: 60/083499
R FILING DATE: 1998-04-29
R APPLICATION NUMBER: 60/083559
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R APPLICATION NUMBER: 60/082704
R APLICATION NUMBER: 60/082797
R FILING DATE: 1998-04-22
R APPLICATION NUMBER: 60/083322
R FILING DATE: 1998-04-28
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                                     APPLICATION NUMBER: 60/077632
FILING DATE: 1998-03-11
APPLICATION NUMBER: 60/077649
FILING DATE: 1998-03-11
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APPLICATION NUMBER: 60/078886
FILING DATE: 1998-03-20
APPLICATION NUMBER: 60/078939
FILING DATE: 1998-03-20
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APPLICATION NUMBER: 60/080107
FILING DATE: 1998-03-31
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FILING DATE: 1998-03-31
APPLICATION NUMBER: 60/080327
FILING DATE: 1998-04-01
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FILING DATE: 1998-04-08
APPLICATION NUMBER: 60/081195
APPLICATION APPLIC
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APPLICATION NUMBER: 60/084366
FILING DATE: 1998-05-05
APPLICATION NUMBER: 60/084414
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APPLICATION NUMBER: 60/079664
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APPLICATION NUMBER: 60/080333
FILING DATE: 1998-04-01
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FILING DATE: 1998-04-08
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APPLICATION NUMBER: 60/081070
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APPLICATION NUMBER: 60/084639
FILING DATE: 1998-05-07
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FILING DATE: 1998-05-22
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APPLICATION NUMBER: 60/079786
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FILING DATE: 1998-05-07
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FILING DATE: 1998-05-22
FILING DATE: 1998-03-10
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PRICR PELICATION NUMBER: 60/08709
PRICR FILING DATE: 1998-05-28
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PRICR FILING DATE: 1998-05-29
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PRICR FILING DATE: 1998-06-30
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Gaps

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Query Match 30.0%; Score 6; DB 13; Length 1089; Best Local Similarity 100.0%; Pred. No. 4.6e+02; Matches 6; Conservative 0; Mismatches 0; Indels

12 WLRRYG 17 |||||| 687 WLRRYG 692

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYBEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430RLC41
CURRENT APPLICATION NUMBER: 10/10/14,581
CURRENT FILING DATE: 2002-06-18
PRIOR APPLICATION NUMBER: 10/052586
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SECRETED AND TRANSMEWBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430R1C50 CURRENT APPLICATION NUMBER: US/10/175,737 CURRENT FILING DATE: 2002-06-19 Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
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es 0; Indels
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100.0%; Pred. No. 4.6
cive 0; Mismatches
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                                                                                                                            ; Sequence 266, Application US/10175737; Publication No. US20030013153A1; GENERAL INFORMATION:
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APPLICATION NUMBER: 60/062250
APPLICATION NUMBER: 60/063120
APPLICATION NUMBER: 60/063120
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Gurney, Austin L.
Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Gurney, Austin L.
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Goddard, Audrey
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                                  687 WLRRYG 692
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Matches 6; Conserv
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12 WLRRYG 17
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US-10-174-581-266
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                                                                                            RESULT 53
US-10-175-737-266
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CURRENT APPLICATION NUMBER: 105/10/176,758
CURRENT FILING DATE: 2002-06-21
                                                                                                                                                                                                                                                                                                                            APPLICANT: Zhang, Zemin Transhembrane POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: SECRETED AND TRANSHEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430R1C42
CURRENY APPLICATION NUMBER: US/10/174,590
CURRENY FILING DATE: 2002-06-18
Prior application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
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30.0%; Score 6; DB 14; Length 1089;
Best Local Similarity 100.0%; Pred. No. 4.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
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US-10-176-758-266
; Sequence 266, Application US/10176758
; Publication No. US20330008353A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
**POTICANT: Chen, Jian
                                                                        Application US/10174590
o. US20030008352A1
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Wood, William I.
Zhang, Zemin
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Smith, Victoria
Watanabe, Colin K.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Pan, James
Smith, Victoria
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Godowski, Paul J.
Gurney, Austin L.
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ORGANISM: Homo Sapien
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CORGANISM: Homo Sapien
US-10-174-590-266
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                                                                                ; Sequence 266, Applic; Publication No. US20; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/078939
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-20
PRIOR PELING DATE: 1998-03-27
PRIOR APPLICATION NUMBER: 60/079786
PRIOR FILING DATE: 1998-03-27
PRIOR APPLICATION NUMBER: 60/080107
PRIOR FILING DATE: 1998-03-31
PRIOR APPLICATION NUMBER: 60/080194
PRIOR FILING DATE: 1998-03-31
PRIOR APPLICATION NUMBER: 60/080134
PRIOR APPLICATION NUMBER: 60/080327
PRIOR APPLICATION NUMBER: 60/080337
PRIOR APPLICATION NUMBER: 60/080337
PRIOR PILING DATE: 1998-04-01
PRIOR FILING DATE: 1998-04-01
PRIOR FILING DATE: 1998-04-01
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R APPLICATION NUMBER: 60/077649
R APPLICATION NUMBER: 60/07886
R APPLICATION NUMBER: 60/078886
R FILING DATE: 1998-03-20
R FILING DATE: 1998-03-20
R FILING DATE: 1998-03-20
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R APPLICATION NUMBER: 60/068017
R FILING DATE: 1997-12-18
R APPLICATION NUMBER: 60/077450
R FILING DATE: 1998-03-10
R APPLICATION NUMBER: 60/077632
                                                                            APPLICATION NUMBER: 60/069335 FILING DATE: 1997-12-11 APPLICATION NUMBER: 60/069425
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APPLICATION NUMBER: 60/063541
FILING DATE: 1997-10-28
APPLICATION NUMBER: 60/063544
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APPLICATION NUMBER: 60/063564
FILING DATE: 1997-10-28
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APPLICATION NUMBER: 60/063870
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APPLICATION NUMBER: 60/065311
FILING DATE: 1997-11-13
FILING DATE: 1997-10-24
APPLICATION NUMBER: 60/063121
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FILING DATE: 1997-10-29
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APPLICATION NUMBER: 60/066466
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APPLICATION NUMBER: 60/066772
FILING DATE: 1997-11-24
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APPLICATION NUMBER: 60/069870
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PRIOR APPLICATION NUMBER: 60/082568
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/082569
PRIOR FILING DATE: 1998-04-21
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APPLICATION NUMBER: 60/082704
FILING DATE: 1998-04-22
APPLICATION NUMBER: 60/082797
FILING DATE: 1998-04-22
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APPLICATION NUMBER: 60/081195
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APPLICATION NUMBER: 60/081838
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PRIOR APPLICATION NUMBER: 60/083428
PRIOR APPLICATION NUMBER: 60/083495
PRIOR APPLICATION NUMBER: 60/083495
PRIOR APPLICATION NUMBER: 60/083496
PRIOR PELING DATE: 1998-04-29
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PRIOR PLILING DATE: 1998-04-29
PRIOR PLILING DATE: 1998-05-05
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PRIOR PELING DATE: 1998-05-07
PRIOR PELING DATE: 1998-05-15
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PRIOR PELING DATE: 1998-06-10
PRIOR PELING DATE: 1998-06-10
PRIOR PELING DATE: 1998-06-10

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APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C93
                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Zhang, Zemin TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430R1C76 CURRENT APPLICATION NUMBER: US/10/176,749 CURRENT FILING DATE: 2002-06-20 Prior application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612 SEQ ID NO 266
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LENGTH: 1089
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                                     US-10-176-749-266

; Sequence 266, Application US/10176749
; Publication No. US20030017542A1
; GENERAL INFORMATION:
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Watanabe, Colin K.
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Godowski, Paul J.
Gurney, Austin L.
Pan, James
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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ORGANISM: Homo Sapien
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Best Local Similarity
Matches 6; Conserv
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CURRENT APPLICATION NUMBER: US/10/176,483
CURRENT FILING DATE: 2002-06-20
Prior application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
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Best Local Similarity 100.0%; Pred. No. 4.6
Matches 6; Conservative 0; Mismatches
                                                        PRIOR APPLICATION NUMBER: 60/088825
PRIOR PAPLICATION NUMBER: 60/088825
PRIOR PELLING DATE: 1998-06-10
PRIOR PELLING DATE: 1998-06-10
PRIOR PELLING DATE: 1998-06-10
PRIOR PELLING DATE: 1998-06-11
PRIOR APPLICATION NUMBER: 60/088863
PRIOR PELLING DATE: 1998-06-11
PRIOR APPLICATION NUMBER: 60/088976
PRIOR PILING DATE: 1998-06-11
PRIOR PELLING DATE: 1998-06-11
PRIOR PILING DATE: 1998-06-12
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FILING DATE: 1998-06-10
APPLICATION UNDRER: 60/088824
FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088825
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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Matches 6; Conserv
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US-10-176-483-266
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LENGTH: 1089
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APPLICANT:

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TITLE OF INVENTION: SCRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACTDS ENCODING THE SAME FILE REPERENCE: P843 ORTIC45
CURRENT APPLICATION NUMBER: 105/10/175,738
CURRENT FILING DATE: 2002-06-19
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C60
CURRENT APPLICATION NUMBER: US/10/175,752
CURRENT APPLICATION NUMBER: US/10/175,752
CURRENT FILING DATE: 2002-06-19
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    Indels
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NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
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Mismatches
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Publication No. US20030022295A1
GRNERAL INFORMATION:
APPLICANT: Baker Kevin P.
APPLICANT: Chen, Jian
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Wood, William I.
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Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
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Best Local Similarity luv...
Section 6; Conservative
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6; Conservative
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Smith, Victoria
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Smith, Victoria
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Goddard, Audrey
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APPLICANT: Chen, Jian
APPLICANT: Desnoyers, Luc
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ORGANISM: Homo Sapien
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US-10-175-752-266
                                                                              687 WLRRYG 692
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US-10-175-738-266
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US-10-175-752-266
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CURRENT APPLICATION NUMBER: US/10/176,915
CURRENT FILING DATE: 2002-06-21
Prior Application removed - See File Wrapper or Palm
SEQ ID NO 266
IBNGTH: 1089
TYPE: PRT
CHARM: Homo Sapien
US-10-176-915-266
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APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Matanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Zhang Zemin
III.
III.E OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
CURRENT APPLICATION NUMBER: US/10/173,706
CURRENT FILING DATE: 2002-06-17
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NUMBER OF SEQ ID NOS: 612
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                                                                                                         Sequence 266, Application US/10176915
Publication No. US20030017544A1
GENERAL INFORMATION:
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Publication No. US20030022293A1
GENERAL INFORMATION:
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Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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Godowski, Paul J.
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Best Local Similarity 100.
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            687 WLRRYG 692
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LENGTH: 1089
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430RIC153
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                                                                     Length 1089;
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CURRENT FILING DATE: 2002-06-25
Prior Application removed - See File Wrapper or Palm
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CURRENT FILING DATE: 2002-06-20
Prior Application removed - See file Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
                                                                     Query Match 30.0%; Score 6; DB 14; Le Best Local Similarity 100.0%; Pred. No. 4.6e+02; Matches 6; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                               ; Sequence 266, Application US/10176913; Publication No. US20030022298A1; GENERAL INFORMATION:
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Watanabe, Colin K.
Wood, William I.
Zhang, Zemin
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Pan, James
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Smith, Victoria
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Goddard, Audrey
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               ; ORGANISM: Homo Sapien
US-10-176-757-266
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US-10-176-913-266
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TYPE: PRT
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CURRENT APPLICATION NUMBER: US/10/176,757
CURRENT FILING DATE: 2002-06-20
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3440R1C70
CURRENT APPLICATION NUMBER: US/10/176,482
CURRENT FILING DATE: 2002-06-20
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                 h Similarity 100.0%; Score 6; DB 14; Length 1089; Similarity 100.0%; Pred. No. 4.6e+02; 6; Conservative 0; Mismatches 0; Indels
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NUMBER OF SEQ ID NOS: 612
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Publication No. US20030022297A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
                                                                                                                                                                                                                               Sequence 266, Application US/10176482
Publication No. US20030022296A1
GENERAL INFORMATION:
APPLICANT: Baker Kevin P.
APPLICANT: Chen, Jian
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Wood, William I.
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Gurney, Austin L.
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Smith, Victoria
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Goddard, Audrey
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US-10-176-482-266
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LENGTH: 1089
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; Sequence 266, Application US/10174572
; Publication No. US20030027263A1
; GENERAL INFORMATION:
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Publication No. US20030027264A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
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Watanabe, Colin K.
Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
                                                                                                                                                          Query Match
Best Local Similarity 100.0
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Goddard, Audrey
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                                                                                                          ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-173-700-266
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US-10-174-572-266
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LENGTH: 1089
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TITLE REPREBRICE: 1934 SORICL47
CURRENT APPLICATION NUMBER: US/10/180,557
CURRENT FILING DATE: 2002-06-25
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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                                                                                                                 Query Match 30.0%; Score 6; DB 14; Length 1089; Best Local Similarity 100.0%; Pred. No. 4.6e+02; Matches 6; Conservative 0; Mismatches 0; Indels
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Wood, William I.
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Gurney, Austin L.
Pan, James
Smith, Victoria
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Wood, William I.
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Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
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; NUMBER OF SEQ ID NOS: 612
; SEQ ID NO 266
; LENGTH: 1089
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Desnoyers, Luc
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                                                               ; ORGANISM: Homo Sapien
US-10-180-552-266
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CORGANISM: Homo Sapien
US-10-180-557-266
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LENGTH: 1089
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APPLICANT: Smith, Victoria
APPLICANT: Matanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Matanabe, Colin K.
APPLICANT: Alang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERBNCE: P3430R1C40
CURRENT APPLICATION NUMBER: US/10/174,572
CURRENT FILING DATE: 2002-06-18
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CURRENT APPLICATION NUMBER: US/10/173,700
CURRENT FILING DATE: 2002-06-17
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
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100.0%; Pred. No. 4.6e+02;
tive 0; Mismatches 0;
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APPLICANT: Wood, William I.
APPLICANT: Mood, William I.
APPLICANT: Zhang, Zemin
TILLE OF INVENTION: SCRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TILLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P34 90R1C28
CURRENT PAPLICATION NUMBER: US/10/174,588
CURRENT FILING DATE: 2002-06-18
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 266
LENGTH: 1089
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APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
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CURRENT APPLICATION NUMBER: US/10/175,739
CURRENT FILING DATE: 2002-06-19
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US-10-175-739-266
i Sequence 266, Application US/10175739
i Publication No. US2033027267A1
i GENERAL INFORMATION:
i APPLICANT: Baker, Kevin P.
i APPLICANT: Chen, Jian
    Watanabe, Colin K.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
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ORGANISM: Homo Sapien
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ORGANISM: Homo Sapien
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Best Local Similarity
Matches 6; Conserv
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: 194430R1031 CURRENT APPLICATION NUMBER: US/10/174,579
CURRENT FILING DATE: 2022-06-18
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
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100.0%; Pred. No. 4.6
tive 0; Mismatches
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Publication No. US20030027265A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Godowski, Paul J.
Gurney, Austin L.
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                                                                                                                                                                                          TYPE: PRT CORGANISM: Homo Sapien US-10-174-579-266
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US-10-174-582-266
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LENGTH: 1089
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LENGIH: 108
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PRIOR FILING DATE: 1997-10-31
PRIOR PRINK DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/066120
PRIOR PRIOR TELLING DATE: 1997-11-13
PRIOR PRIOR PRIOR DATE: 1997-11-21
PRIOR PRIOR PRIOR DATE: 1997-11-24
PRIOR PRIOR APPLICATION NUMBER: 60/06672
PRIOR PELING DATE: 1997-11-24
PRIOR PELING DATE: 1997-11-12
PRIOR PELING DATE: 1997-11-12
PRIOR PELING DATE: 1997-12-18
PRIOR PELING DATE: 1998-03-20
PRIOR PELICATION NUMBER: 60/08139
PRIOR PELICATION NUMBER: 60/08139
PRIOR PELICATION NUMBER: 60/081499
PRIOR PELICATION NUMBER: 60/081499
PRIOR PELICATION NUMBER: 60/081499
PRIOR PELICATION NUMBER: 60/081499
PRIOR PELING DATE: 1998-04-29
PRIOR PELICATION NUMBER: 60/081499
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PRIOR PELICATION NUMBER: 60/081399
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APPLICANT: ADDISONAL COLLINE.

APPLICANT: Zhang, Zemin

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC

TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C61

CURRENT APPLICATION NUMBER: US/10/175,740

CURRENT FILING DATE: 2002-06-18

Prior Application removed - See File Wrapper or Palm

SEQ ID NO 266

LENGTH: 1089
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APPLICANT: Waterabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Thang, Zemin I.
APPLICANT: Thang, Zemin I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
TILLS EFERENCE: P3439RICS2
CURRENT APPLICATION NUMBER: U5/10/175,743
CURRENT PILING DATE: 2002-06-16
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Similarity 100.0%; Pred. No. 4.6
6; Conservative 0; Mismatches
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US-10-175-743-266
; Sequence 266, Application US/10175743
; Publication No. US2030027269A1
; GENERAL INFORMATION:
; APPLICANT: Baker Kevin P.
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PRIOR FILING DATE: 2002-01-15
PRIOR PAPPLICATION NUMBER: 60/059263
PRIOR PELLING DATE: 1997-09-18
PRIOR PELLING DATE: 1997-09-18
PRIOR PAPLICATION NUMBER: 60/069266
PRIOR PELLING DATE: 1997-09-18
PRIOR PELLING DATE: 1997-09-18
PRIOR PELLING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063120
PRIOR PELLING DATE: 1997-10-24
PRIOR PELLING DATE: 1997-10-21
PRIOR PELLING DATE: 1997-10-21
PRIOR PELLING DATE: 1997-10-21
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PRIOR APPLICATION NUMBER: 60/063541
PRIOR FILING DATE: 1997-10-28
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PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063564
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PRIOR APPLICATION NUMBER: 60/063734
PRIOR FILING DATE: 1997-10-29
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APPLICATION NUMBER: 60/063564
                         Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
Gurney, Austin L.
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Godowski, Paul J
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Smith, Victoria
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Best Local Similarity
Matches 6; Conserv
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PRIOR APPLICATION NUMBER: 60/084640
PRIOR APPLICATION NUMBER: 60/084641
PRIOR APPLICATION NUMBER: 60/08573
PRIOR PILING DATE: 1938-05-15
PRIOR PAPLICATION NUMBER: 60/08520
PRIOR APPLICATION NUMBER: 60/08623
PRIOR APPLICATION NUMBER: 60/08623
PRIOR APPLICATION NUMBER: 60/086392
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PRIOR APPLICATION NUMBER: 60/086292
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PRIOR APPLICATION NUMBER: 60/086292
PRIOR PILING DATE: 1938-05-22
PRIOR APPLICATION NUMBER: 60/086292
PRIOR PILING DATE: 1938-05-20
PRIOR PILING DATE: 1938-05-20
PRIOR PILING DATE: 1938-05-20
PRIOR PILING DATE: 1938-05-20
PRIOR PILING DATE: 1938-05-06-03
PRIOR PILING DATE: 1938-05-06-03
PRIOR PILING DATE: 1938-05-06-03
PRIOR PILING DATE: 1938-06-03
PRIOR PILING DATE: 1938-06-03
PRIOR PAPLICATION NUMBER: 60/08029
PRIOR PAPLICATION NUMBER: 60/08029
PRIOR PILING DATE: 1938-06-03
PRIOR PAPLICATION NUMBER: 60/08029
PRIOR PAPLICATION NUMBER: 60/08029
PRIOR PILING DATE: 1938-06-05
PRIOR PILING DATE: 1938-06-10
PRIOR PILING DATE
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430R1C119
CURRENT APPLICATION NUMBER: US/10/176,488
CURRENT FILING DATE: 2002-06-21
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Pred. No.
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Pred. No.
PRIOR APPLICATION NUMBER: 60/089105
PRIOR FILING DATE: 1998-06-12
PRIOR APPLICATION NUMBER: 60/089512
PRIOR APPLICATION NUMBER: 60/089514
PRIOR PLING DATE: 1998-06-16
PRIOR PLING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/089538
PRIOR FILING DATE: 1998-06-17
PRIOR FILING DATE: 1998-06-17
PRIOR FILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089598
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Publication No. US20030027271A1
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100.0%; Pre
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Smith,Victoria
Watanabe,Colin K.
Wood,William I.
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Gurney, Austin L.
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Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity 100.
Matches 6; Conservative
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CRGANISM: Homo Sapien
US-10-176-488-266
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